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Oncogene 1999 Jan 14	:18(2):559-64	

A tumor specific single chain antibody dependent gene expression system.

Mary MN, Venot C, Caron de Fromentel C, Debussche L, Conseiller E, Cochet O, Gruel N, Teillaud JL, Schweighoffer F, Tocque B, Bracco L

Gene Medicine Department, Rhone-Poulenc Rorer S.A., Vitry-sur-Seine, France.

The design of conditional gene expression systems restricted to given tissues or cellular types is an important issue of gene therapy. Systems based on the targeting of molecules characteristic of the pathological state of tissues would be of interest. We have developed a synthetic transcription factor by fusing a single chain antibody (scFv) directed against p53 with the bacterial tetracycline repressor as a DNA binding domain. This hybrid protein binds to p53 and can interact with a synthetic promoter containing tetracycline-operator sequences. Gene expression can now be specifically achieved in tumor cells harboring an endogenous mutant p53 but not in a wild-type p53 containing tumor cell line or in a non-transformed cell line. Thus, a functional transactivator centered on single chain antibodies can be expressed intracellularly and induce gene expression in a scFv-mediated specific manner. This novel class of transcriptional transactivators could be referred as 'trabodies' for transcription-activating-antibodies. The trabodies technology could be useful to any cell type in which a disease related protein could be the target of specific antibodies.

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Oncogene 1999 Jan 14;18(2):551-7

Restoration of transcriptional activity of p53 mutants in human tumour cells by intracellular expression of anti-p53 single chain Fv fragments.

Caron de Fromentel C, Gruel N, Venot C, Debussche L, Conseiller E, Dureuil C, Teillaud JL, Tocque B, Bracco L

Gene Medicine Department, Rhone-Poulenc Rorer SA, Vitry sur Seine, France.

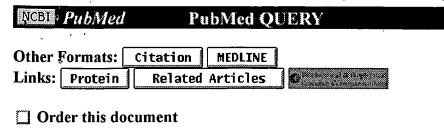
We report here the production and the properties of single chain Fv fragments (scFvs) derived from the anti-p53 monoclonal antibodies PAb421 and 11D3. 11D3 is a newly generated monoclonal antibody which exhibits properties very comparable to those of PAb421. The scFvs PAb421 and 11D3 are able to stably associate with p53 and to restore the DNA binding activity of some p53 mutants in vitro. When expressed in p53 -/-human tumour cells, the scFv421 is essentially localized in the cytoplasm in the absence of p53, and in the nucleus when exogenous p53 is present. Thus, p53 is also able to stably associate with an anti-p53 scFv in cells. Cotransfection of p53 -/- human tumour cells with expression vectors encoding the His273 p53 mutant and either scFv leads to restoration of the p53 mutant deficient transcriptional activity. These data demonstrate that, in human tumour cells, these scFvs are able to restore a function essential for the tumour suppressor activity of p53 and may represent a novel class of molecules for p53-based cancer therapy.

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PubMed



Biochem Biophys Res Commun 1997 Jan 13;230(2):242-6

Characterization of scFv-421, a single-chain antibody targeted to p53.

Jannot CB, Hynes NE

Friedrich Miescher Institute, Basel, Switzerland.

A gene encoding a single-chain antibody (scFv) which specifically binds the tumor suppressor protein p53 has been constructed from RNA of hybridoma cells producing Pab 421. scFv-421 which was expressed and purified from bacteria specifically binds p53. scFv-421, as well as the previously described scFv-FRP5 and -R1R (1), were expressed intracellularly in mammalian cells and targeted to different subcellular locations, including the nucleus, cytoplasm, and endoplasmic reticulum (ER). High levels of all ER targeted scFv proteins, but not nuclear or cytoplasmic targeted proteins, were found in transfected COS-1 cells. In an attempt to stabilize the proteins, sequences encoding the mouse immunoglobin CK constant domain were added to each scFv construct. This led to a moderate increase in the cytoplasmic expression of scFv-FRP5.

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Oncogene 1998 Nov 12;17(19):2445-56

Characterization of a new intrabody directed against the N-terminal region of human p53.

Cohen PA, Mani JC, Lane DP

Department of Biochemistry, Medical Sciences Institute, University of Dundee, UK.

Genes encoding the rearranged immunoglobulin heavy and light chain variable regions of DO-1, a monoclonal antibody directed against human p53, have been used to construct a single-chain antibody. DO-1 recognizes an N-terminal epitope in the region involved in the transactivation function of p53 and the binding of Mdm2. The DO-1 single chain scFv expressed in the periplasm of E. coli or at the surface of the filamentous phage M13 retained the immunological specificity and affinity of the full length antibody. Furthermore, the DO-1 recombinant antibody was able to inhibit the in vitro binding of Hdm2, and was shown to be a powerful protecting agent of p53's DNA binding activity at 37 degrees C. The DO-1 single-chain antibody has been used to construct single-chain intracellular antibodies (intrabodies) for expression in the cytoplasm and the nucleus of mammalian cells. These anti-p53 intrabodies were additionally modified by addition of a Ckappa domain to increase cytoplasmic and nuclear stability. Here we show that expression of the DO-1 single-chain antibody in the H1299 cell line results in an inhibition of p53's transactivation function. The DO-1 intrabody is a useful tool to study those functions of p53 driven by the N-terminal region of the protein.

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Restoring p53-dependent trans-activating activity to cell containing mutant p53 - by delivering single-chain antibody specific for the

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Pred. No. 2.6e-88;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                          Conjugate; cell targeting; cytotoxic drug;
cell surface antigen; treatment; cancer; ir
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N-PSDB; V72064.
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.4e-67;
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                                                                   Single chain antibody; ScFv 421; mouse; p53 protein; oligomerisation; regulatory domain; p53 mutant; H273; W248; G281; p53-dependent trans-activating activity; restoration; tumour-suppressing activity; tumour cell; treatment; hyper-proliferation; cancer; re-stenosis; ss.
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Pred. No. 8.5e
21; Mismatches
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.5e-67;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Restoring p53-dependent trans-activating activity to cell mutant p53 - by delivering single-chain antibody specific mutant, particularly for treatment of tumours
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1998-272140/24
                                                                                                                                                                                                                                  FGGGTKLEIK 249
                                                                                                                                                                                                                                                                           QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
                                                                                                                                                                                                                                                                                                                                  SSGGGGSGGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLENSRTRKNYLAWYQ
                                                                                                                                                                                                                                                                                                                                                                          apkfqgkatmtadtssntaylqlsslasedtavyycn----fygdaldywgqgttvtv
                                                                                                                                                                                                                                                              qrpgqspkrliylvskldsgvpdrftgsgsgtdftlkinrveaedlgvyycwqgthsplt
                                                                                                                                                                                                                                                                                                                  ssggggsgggsggsdvlmtqtpltlsvtigqpasisckssqsllds-dgktylnwll
                                                                                                                                                                                                                                                                                                                                                                                                       APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV
                                                                                                                                                                                                                                                                                                                                                                                                                                 qvqlqqsgaelvrsgasvklsctasgfnikdyymhwvkqrpeqglewigwidpengdtey
                                                                                                                                                                                                        fgagtkleik 243
                                                   p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            187;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 31;
                                                 protein variant S-325 encoded by pEC176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                              (first
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                                                                              entry)
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74.8%;
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                                                                                                                                      535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 961; DB 19;
Pred. No. 1.1e-62;
3; Mismatches 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                         239
                                                                                                                                                                                                                                                                                                                                                                             114
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Leucine zipper o substitution; re anti-oncogene; i

r domain; LZD; oligomerisation domain; mutant replacement; transactivation; viral protein ; oyperproliferation; cancer; restenosis; Scf

mutant;

mutein

Human

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                                                                                                                                                                                                                                                                                                                                                                                                   Claimed variants of protein p53 have at least part of the coligomerisation domain deleted and replaced by a leucine zipper comain. The mutants preferably also have at least part of the p53 transactivation domain (amino acids 1-74) deleted and replaced by the transactivating domain (TD) from herpes simplex virus viral protein VP16 (amino acids 411-490) or by a protein domain able to bind selectively to a transactivator, especially a single-chain comparison are apecifically claimed p53 variant designated s-325 and comprising a specifically claimed p53 variant designated s-325 and comprising a ScFv domain, amino acids 75-325 of human wild-type p53 and a core active and more stable tumour suppressors and apoptosis-inducing agents than wild-type p53 and are active where the wild-type protein cis not, i.e. they are not inactivated by dominant negative or oncogenic mutants, nor by other cellular proteins (because the leucine zipper comain prevents formation of inactive mixed oligomers).
                                                                                                                                                                                                                                                                                                                    Query Mat
Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chimeric -
Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 36; Pages 88-90; 133pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New p53 variants e.g. with oligomerisation domain replaced by leucine zipper - useful for treating hyper-proliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-132633/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9704092-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumour suppression;
                                                                176
236
                                240
                                                                                                                              117
                                                                                                                                                          121
                                                                                                                                                                                          63
                                                                                                                                                                                                                         61
                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                       1 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60
                                                                                                                                                                                                                                                                                                                                                      Match
                                                                                                                                                                                                         APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYMGQGTTVTV 120
                                FGGGTKLEIK 249
                                                           qrpgqspkrliylvskldsgvpdrftgsgsgtdftlkinrveaedlgvyycwqgthsplt
                                                                                             QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
                                                                                                                           ssggggsgggggggggdvlmtqtpltlsvtigqpasisckssqsllds-dgktylnwll
                                                                                                                                            SSGGGGSGGGGSGGGSDIELTOSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                        apkfqgkatmtadtssntaylqlsslasedtavyycn----fygdaldywgqgttvtv
                                                                                                                                                                                                                                                     qvqlqesgaelvgsgasvklsctasgfnikdyymhwvkqrpeqglewigwldpengdtey 62
   fgagtklelk
                                                                                                                                                                                                                                                                                                                       al Similarity
186; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
Herpes simplex virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conseiller
                                                                                                                                                                                                                                                                                                                                                                                                   535 AA
                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95FR-0008729
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   245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     apoptosis; single chain antibody variable domain.
                                                                                                                                                                                                                                                                                                                                    72.6%;
                                                                                                                                                                                                                                                                                                                       23;
                                                                                                                                                                                                                                                                                                                    Score 955; DB
Pred. No. 7.2e
23; Mismatches
                                                                                                                                                                                                                                                                                                                     DB 18;
7.2e-62;
nes 33;
                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                       Indels
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                                                                235
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W28492
ID W
Query Match
Best Local S
Matches 186
                                                                                                                                                the transactivating domain (TD) from herpes simple virus viral protein VPI6 (amino acids 411-490) or by a protein domain able to bind selectively to a transactivator, especially a single-chain antibody variable domain (ScFv). The present sequence is that of a specifically claimed p53 variant designated S-325H and comprising a ScFv domain, amino acids 75-325 of human wild-type p53 (but with Arg182 replaced by His) and a leucine zipper domain at the C-terminal. The p53 variants are more active and more stable tumour suppressors and appropriate insuring agents than wild-type p53 and are active where the wild-type protein is not, i.e. they are not inactivated by dominant negative or oncogenic mutants, nor by other cellular proteins (because the leucine zipper domain prevents formation of inactive mixed
                                                                                                                                                                                                                                                                                                                                              Claimed variants of protein p53 have at least part of the oligomerisation domain deleted and replaced by a leucine zipper domain. The mutants preferably also have at least part of the p53 transactivation domain (amino acids 1-74) deleted and replaced by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Leucine zipper domain; LZD; oligomerisation domain; mutant; mutein; substitution; replacement; transactivation; viral protein VP16; HSV; anti-oncogene; hyperproliferation; cancer; restenosis; ScFv; tumour suppression; apoptosis; single chain antibody variable domain
                                                                             Sequence
                                                                                                            been
                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New p53 variants e.g. with oligomerisation domain replaced by leucine zipper - useful for treating hyper-proliferative disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric -
                                                                                                                                        oligomers).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bracco L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W28492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W28492 standard; Protein; 535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RHON ) RHONE POULENC RORER SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9704092-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-NOV-1997
                                                                                                         Note: this sequence does not appear in the specification and has seen produced by modifying the given sequence of variant V-325).
                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1997-132633/12.
                                                                                                                                                                                                                                                                                                                                                                                                                          36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p53 protein variant S-325H.
                                                                                                                                                                                                                                                                                                                                                                                                                          Page -; 133pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conseiller
                                                                             535 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Herpes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         and restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95FR-0008729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96WO-FR01111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Arg residue at position 182 of wild-type
p53 has been mutated to His"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     simplex virus
                72.68;
   23;
Score 955; DB
Pred. No. 7.2e
23; Mismatches
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                .2e
   18;
-62;
33;
                              Length
   Indels
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        30-DEC-1998
                                                                               Misc-difference
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                                                                                                                                    Region
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                                                                                                                                                                                                                                                                                                              Key
                                                                                                                                                                                                                                                                                                                                Chimeric
                                                                                                                                                                                                                                                                                                                                        Chimeric
                                                                                                                                                                                                                                                                                                                                                 Chimeric -
                                                                                                                                                                                                                                                                                                                                                                   allergy; therapy;
BiTAbB7-24-1G10H6.
                                                                                                                                                                                                                                                                                                                                                                          B7 binding molecule; costimulatory molecule; B7.1; CD80; B7.2; CD86; T cell activation; inhibitor; graft versus host disease; transplant rejection; autoimmune disease; allergy; therapy; human; bispecific tetravalent antibody; BiTAb;
                                                                                                                                                                                                                                                                                                                                                                                                                        Bispecific tetravalent antibody BiTAbB7-24-IG10H6
                                                                                                                                                                                                                                                                                                                                                                                                                                           10-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                             W90217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W90217 standard; Protein; 580 AA.
                          WO9858965-A2
                                                                                                                                                                                                                                                                Peptide
                                                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                    Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APNFQGKATTTADTSSNTAYLHLSSLTSEDTTYYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     apkfqgkatmtadtssntaylqlsslasedtavyycn-----fygdaldywgqgttvtv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fgagtklelk 245
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                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                 Mus sp.
                                                                                                                                                                                                                                                                                                                                synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                               /note= ", 322..327
                                                                                                                                    /note= "
462..574
                                                                                                                                                                                                                                                                                                                                         sapiens
                                                                                                                                                                                                            /note=
274..3
                                                                                                                                                      447..461
                                                                                                                                                                               /note=
                                                                                                                                                                                                                             263..273
                                                                                                                                                                                                                                                                                    25..138
                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                           /note-
                                                                                                                                                               /note=
                                                                                                                                                                                                                                       /note-
                                                                                                                                                                                                                                                154..262
                                                                                                                                                                                                                                                                  139..153
                                                                                                                                                                                                                                                                                          /note= "pelB signal peptide"
                                                                                                          'note=
                                                                                                                                                                                                  'note=
                                                                                                                                                                                                                                                        note-
                                                                                                                                                                                                            ..308
                                                                                                                                                                         . . 446
                                                                                                                                                                                          ..319
                                                                                      "encoded by
                                                                                                                           "VL region
                                                                                                                                             "(G4S3)flexible linker"
                                                                                                                                                                                                                                                        "(G4S3) flexible linker"
                                                                                                                                                                                                                                                                          "VH region anti B7.1 MAb"
                                          "codons for these amino acids are not
present in the DNA sequence for
BTITADB7-24-1G1-H6 provided in the
specification"
                                                                                                                                                                                                  "helix-turn-helix dimerisation
                                                                                                          "His6
                                                                                                                                                              "VH region anti B7.2 MAb
                                                                                                                                                                               "human IgG3
                                                                                                                                                                                                                   "human IgG3 hinge region"
                                                                                                                                                                                                                                     "VL region
                                                                                                                            anti B7.2 MAb
                                                                                                                                                                                                                                      anti B7.1 MAb'
                                                                                       CIG.
                                                                                                                                                                                hinge domain"
                                                                                                                                                                                                 domain"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       XXXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BITAb is a homodimer of 2 identical molecules, each containing both (an anti B7.1 and anti B7.2 scFv (bispecificity). An anti-B7.1 and anti-B7.2 scFv are linked using a dimerisation domain (see W90219), which drives the homodimerisation of the molecule. DNA (see X01651) encoding the BITAb has been constructed to allow expression of the BITAb in transformed E. coli cells. The BITAb cross-links, and/or cross-reacts, with the costimulatory molecules B7.1 and B7.2 that are expressed on the membrane of professional antigen-presenting cells, leading to the inhibition of antigen-cross-reacts antigen-presenting cells, reading to the inhibition of antigen-cross-collection. The invention relates to such B7-binding molecules, methods for their production, and their use for treating or preventing diseases of the immune system, in particular graft rejection, graft versus host disease, allergy and continuous diseases (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                            W90218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This polypeptide comprises the bispecific tetravalent antibody BiTAbB7-24-IG10H6. The molecule consists of 4 scFvs, i.e. 2 anti B7.1 scFvs and 2 anti B7.2 scFvs (tetravalency). One single BiTAb is a homodimer of 2 identical molecules, each containing botl an anti B7.1 and anti B7.2 scFv (bispecificity). An anti-B7.1 and
Bispecific
                                   10-MAY-1999
                                                                                                     W90218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 7.1; Fig. 16; 182pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New molecules which bind B7.1 and B7.2 - useful to immune diseases including allograft rejection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bosman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INNO-) INNOGENETICS NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-JUN-1998;
                                                                                                                                                                                       565
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                                                                                                                                                                                                                                                                                            181
                                                                                                                                                                                                                                                                                                                            445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   325
                                                                                                                                                                                                        241 GGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                          385 npkfegkatltvdkssstaymqlesltsedsavyycarfayygdyyyimdywgqgttvtv
                                                                                                                                                                                                                                                                                                                                                              121
                                                                                                                                                                                                                                                                                                                                                                                                                                 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
nes 183; Conserv
                                                                                                                                     8
                                                                                                                                                                                                                                                                            QRPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLPTF
                                                                                                                                                                                     gggtkleik
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                                                                                                                                                                                                                                                                                                                                          SSGGGGSGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                                                                                        ssggggsggggggggdieltqspsslavsageevtmtckssqsvlyssnqknylawyq
                                                                                                                                                                                                                                                                                                                                                                                                                                APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                qvqlqqsgpelekpgasvkisckasgysftghnmnwvkqsngkslewigiidpyyggtsy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
                                                                                                     standard;
tetravalent antibody BiTAb1G10-B7-24H6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Buyse M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       580 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                 (first entry)
                                                                                                                                                                                       573
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                                                                                                   Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 72.3%; Score 951; D)
73.5%; Pred. No. 1.50
tive 22; Mismatches
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B7 binding molecule; costimulatory molecule; B7.1; CD80;

в7.2;

73.3%;

6.5e-61;

0

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BITAbİGİÖ-B7-24H6. The molecule consists of 4 scFvs, i.e. 2 anti B7.1 scFvs and 2 anti B7.2 scFvs (tetravalency). One single BITAb is a homodimer of 2 identical molecules, each containing both an anti B7.1 and anti B7.2 scFv (bispecificity). An anti-B7.1 and and anti-B7.2 scFv are linked using a dimerisation domain (see w90219), which drives the homodimerisation of the molecule. DNA (see x01652) encoding the BiTAb has been constructed to allow expression of the BiTAb in transformed E. coli cells. The BiTAb cross-links, and/or cross-reacts, with the costimulatory molecules B7.1 and B7.2 that are expressed on the membrane of professional antigen-presenting cells, leading to the inhibition of antigen-specific T cell activation. The invention relates to such
Seguence
                        antigen-presenting cells, leading to the inhibition of an specific T cell activation. The invention relates to such B7-binding molecules, methods for production, and tifor treating or preventing diseases of the immune system, particular graft rejection, graft versus host disease, all autoimmune diseases (claimed)
                                                                                                                                                                                                                          This polypeptide comprises the bispecific tetravalent BiTAblGlO-B7-24H6. The molecule consists of 4 scFvs,
                                                                                                                                                                                                                                                                                        New molecules which bind {\sf B7.1} and {\sf B7.2} - useful to immune diseases including allograft rejection
                                                                                                                                                                                                                                                                                                                                                                      Bosman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CD86; T cell activation; inhibitor; graft versus host disease; transplant rejection; allograft rejection; autoimmune disease; allergy; therapy; human; bispecific tetravalent antibody; BiTAb; BiTAb1G10-B7-24H6.
                                                                                                                                                                                                                                                                Example 7.1;
                                                                                                                                                                                                                                                                                                                                                                                                (INNO-) INNOGENETICS
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260..285
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427..441
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note= "(G4S3) flexible linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..426
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Query Match

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      This is the amino acid sequence of a fusion protein comrising of agric single chain antibody (with the heavy and light chain joing a linker), fused to the clavin protein. Clavin is an inhibitor protein, and functions by inactivating the ribosomes. Clavin or conjugates are useful as anticancer and/or antiviral agents. The
                                                                    Claim
                                                                                                 DNA encoding Aspergillus protein, clavin - useful
                                                                                                                              N-PSDB;
                                                                                                                                                            Colnaghi M,
                                                                                                                                                                                                                                                                                                                                               Chimeric
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                                                                                                                                                                                                                                                                 W09749726-A1.
                                                                                                                                                                                                                                                                                             Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                         Nucleotide sequence encoding the Mgr6-clavin immunotoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTVSS 122
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                                                                                                                                                             Mele
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The invention relates to a method of identifying binding site domains (BSD) that retain the capacity of binding to a predetermined epitope when positioned C-terminal of at least one further domain in a recombinant bior multivalent polypeptide. The method comprises (a) testing a panel of a BSD displayed on the surface of a biological display system as part of a fusion protein for binding to a predetermined epitope, where the fusion protein comprises an additional domain positioned N-terminal of the BSD of the BSD displayed on the BSD displayed on the SSD displayed on the
                                                                                                                                                                                                                                             Phage display system for identification of binding
                                                                                                                                                                                                                                                                                                             WPI; 1999-338004/28.
                                                                                                                                                                                                                                                                                                                                                      Borschert K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9925818-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Binding site domain; autoimmune disease; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse scFV
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                                                                                                                                                                             Claim 27;
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                                                                                                                                                                             Fig 6.6; 152pp; English.
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scFv-antibody;
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Pred. No. 2.4e-60;
5; Mismatches 36
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single-chain Fv; mouse.
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RESULT 11
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       Phage display system for identification retaining capacity to bind an epitope
                                               WPI; 1999-338004/28
N-PSDB; X77247.
                                                                                    Borschert K,
                                                                                                                                       17-NOV-1997;
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                                                                                                                                                                                                                                                                  autoimmune
                                                                                                                                                                                                                                                                                                      Mouse scFV fragment 5-10.
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                                                                                                        (KUFE/) KUFER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             scFV fragments.
                                                                                                                                                                                                                                                          site domain; BSD; epitope;
une diseáse; scFv-antibody;
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                                                                                    Kufer
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                                                                                                                                                               98WO-EP07313
                                                                                    Ρ,
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72.9%; Pred. No. 2e-60;
tive 25; Mismatches
                                                                                    Lutterbuese
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single-chain Fv; mouse.
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                      of.
                                                                                    Raum
                     binding site
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                      domains
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Claim 27; Fig 6.10; 152pp; English

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Matches 177;
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                                                                                                                                                                                                                                                                                                                                                                Conjugate; cell targeting; cytotoxic drug; plasmid; fusion protein;
prodrug-converting enzyme; cell surface antigen; treatment; cancer;
                               10-MAY-1997;
                                                                                        05-MAY-1998;
                                                                                                                                                   19-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fusion protein
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                                                                                                                                                                                                                                                                                                                                    inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gtkleik 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTVSSG 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        legsgaelvrpgtsvkisckasgyaftnywlgwvkgrpghglewigdifpgsgnihynek 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      248
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                             rheumatoid arthritis; antibody; prodrug therapy system
                                                                                                                                                                                                                                                                                                                                                                                                                                                        pICI266/8061scFvtag/his.
                         97GB-0009421
                                                                                           98WO-GB01294
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71.7%;
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Pred. No. 9e-60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 20; Length 248;
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19-NOV-1998

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RESULT 13
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Best Local S
Matches 173
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                                                                                                               Conjugate; cell targeting; cytotoxic drug; plasmid; fusion protein;
prodrug-converting enzyme; cell surface antigen; treatment; cancer;
                                                                      Synthetic
                                                                                                                                                         Fusion protein (806.077 scFv-CPG2)2
                                                                                                                                                                                        10-MAY-1999
                                                                                                                                                                                                                                                W82745 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New gene construct expressing conjugate of targeting agent and prodrug-converting enzyme - useful for, e.g. targeted production of cytotoxic drug in vivo, especially for treatment of cancer.
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N-PSDB; V72069.
                                         WO9851787-A2
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                                                                                                 inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ZENE ) ZENECA LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
                                                                                                                                                                                                                                                                                                                                       FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                            QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
                                                                                                                                                                                                                                                                                                                                                                                                                                    ssggggsgggcsggggsgivltqspaimsaspgekvtitcsasssv-----tymhwfq
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLENSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fgagtkleik 263
                                                                                                                                                                                                                                                                                                                                                                                qkpgtspklwiystsnlasgvparfsgsgsgtsysltisrmeaedaatyycqqrstyplt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         281 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                      (first entry)
                                                                                               rheumatoid arthritis; antibody; prodrug therapy system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68.9%; Score 906; DB 20; 69.2%; Pred. No. 1.3e-58;
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Y17965
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence is a used in a method for obtaining a novel gene construct (A) which expresses, in cells of a mammal, a conjugate (B) of a cell-targeting group (I) and a heterologous prodrug-converting enzyme (II), and (B) is directed to leave the cell for selective localisation at a cell surface antigen (Ag) recognised by (I). Delivery of (A) to a target site, then administration of (III) is used for targeted release of cytotoxic drug, specifically for treating cancer but also inflammation such as rheumatoid arthritis. In situ generation of the targeting antibody increases selectivity, reducing side effects at normal tissue. The method is applicable to any antibody-directed enzyme prodrug therapy
WO9925818-A1.
                                                Binding site domain; BSD; epitope; fusion protein; therapeutic; autoimmune disease; scFv-antibody; single-chain Fv; mouse.
                                                                                                                    04-AUG-1999
                                                                                                                                               Y17965
                                                                                                                                                                        Y17965 standard; Protein; 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 17; Page 87-89; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New gene construct expressing conjugate of targeting agent and prodrug-converting enzyme - useful for, e.g. targeted production cytotoxic drug in vivo, especially for treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Blakey DC, Emery SC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAY-1998;
                                                                                          Mouse scFV
                                                                                                                                                                                                                                                                   240
                                                                                                                                                                                                                                                                                           191
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                                                                                                                                                                                                                                                                                                       QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
                                                                                                                                                                                                                                                                                                                                                            SSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKYAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                         fgagtkleik 260
                                                                                                                                                                                                                                                                FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                           qkpgtspklwiystsnlasgvparfsgsgsgtsysltisrmeaedaatyycqqrstyplt
                                                                                                                                                                                                                                                                                                                                                ssggggsgggggggivltqspaimsaspgekvtitcsasssv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           666
                                                                                          fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        68.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 906; DB 20;
Pred. No. 3.2e-58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 666;
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                                                              cancer;
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26-JAN-1996 (first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of identifying binding site domains (CC (BSD)) that retain the capacity of binding to a predetermined epitope when positioned C-terminal of at least one further domain in a recombinant bicompositioned C-terminal of at least one further domain in a recombinant bicom positioned C-terminal of at least one further domain in a recombinant bicom multivalent polypeptide. The method comprises (a) testing a panel of twistion protein for binding to a predetermined epitope, where the fusion protein comprises an additional domain positioned N-terminal of the BSD and an amino acid sequence that mediates anchoring of the fusion protein comparises of the display system; and (b) identifying a BSD that the surface of the display system; and (b) identifying a BSD that comprise antibody binding sites capable of efficiently binding to the comprise antibody binding sites capable cof efficiently binding to the method are useful therapeutically and diagnostically, for e.g. cancer and autoimmune diseases. ScFv-antibody corrections using the new in vitro method. Sequences X17957-965 represent course acre for the method. Sequences X17957-965 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 175;
R75719 standard; Protein; 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Phage display system for identification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-338004/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mouse scFV fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            retaining
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                                                                                   242
                                                                                                          243 GTKLEIK 249
                                                                                                                                               122
                                                                                                                                                                                                                                      124 GGGSGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQQKP 183
                                                                                                                                                                                                                                                                                          65
                                 15
                                                                                                                                                                                                                                                                                                                          64 FQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTVSSG 123
                                                                                                                                                                                                                                                                                                                                                           σ
                                                                                                                                                                                                                                                                                                                                                                           4 LQESGAELVRSGASVNLSCTASGENIKDYYMHWVKQRPEEGLEWIGYIDPESGETEYAPN 63
                                                                                 gtkleik 248
                                                                                                                                                                                                                      gggsggggggggselvmtqspsslamsvgqkvtmsckssqsllnssnqknylawyqqkq
                                                                                                                                                                                                                                                                                                                                                      legsgaelvrogtsvkisckasgyaftnywlgwvkgrpghglewvgdifpgsgnahynek 64
                                                                                                                                                                                                                                                                                      fkgkatltadkssytaymqlssltsedsavyfcarlrnwd---eamdywgqgttvtvssg 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          capacity to bind an epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              248 AA;..
                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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70.9%;
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%; Pred. No. 2.5e
27; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 20;
?.5e-58;
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MFE-23 antibody.

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This sequence represents the MFE-23 antibody, which is an antibody against carcinoembryonic antigen (CEA). CEA is a marker antigen for cancer imaging and therapy. The MFE-23 antibody sequence was obtained using phage technology. In this process, mice were immunised with CEA. The antibody variable region genes obtained from these mice were then amplified from cDNA and cloned as a single chain FV (scFV) into bacteriophage vectors, producing a library. The phages that bound to
                                                                                     Antibody for carcinoembryonic antigen of colorectal cancer
                                                                                                               N-PSDB; Q90663.
                                                                                                                                                                         03-DEC-1993;
                                                                                                                       WPI; 1995-215234/28
                                                                                                                                       Begent RHJ,
                                                                                                                                                                                         05-DEC-1994;
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                                                                                                                                                       (CANC-) CANCER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy; diabody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody; MFE-23; carcinoembryonic antigen; CEA; colorectal tumour;
                                                                     ω,
                                                                    Page 48-49; 72pp; English.
                                                                                                                                       Chester
                                                                                                                                                        RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                        93GB-0024807
                                                                                                                                                                                         94WO-GB02658
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27. 1.7
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62..75
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52..61
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162..267
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Search completed: February 12, Job time: 38 sec

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Best Local S
Matches 171
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                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  antitumour agent or a
                                                                                                                                                                                                                                                                                                                                                                                                                                   fragments which bind to two different antigens), and may be linked to antitumour agent or a detectable label.
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                                                                                 sypltfgagtklelk
                                 NLP-TFGGGTKLEIK 249
                                                               mhwfqqkpgtspklwiystsnlasgvparfsgsgsgtsyslt1srmeaedaatyycqqrs
                                                                                                                            ttvtvssggggsggggsggggsenvltqspaimsaspgekvtitcsasssv-----sy
                                                                                                                                         TTVTVSSGGGGSGGGGGGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLENSRTRKNY
                                                                                                                                                                                           apkfqgkatfttdtssntaylqlssltsedtavyycn----egtptgpyyfdywgqg
                                                                                                                                                                                                            APNFQGKATVTADTSSNTAYLHLSSLTSEDTTYYYCNAVIYYEYDG-----YALDYWGQG
                                                                                                                                                                                                                                                       171;
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                                                                                                                                                                                                                                                                                                                                                                                                    270 AA;
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                                                                                                                                                                                                                                                                                                                                   67.18;
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Pred. No. 5.
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                      Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein search, using sw model
                                      100.0
73.1
72.6
                                                                                                                    Match Length DB
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_6/ptodata/2/paa/PCT_NEW_COMB.pep:*
_6/ptodata/2/paa/US05_NEW_COMB.pep:*
_6/ptodata/2/paa/US07_NEW_COMB.pep:*
_6/ptodata/2/paa/US08_NEW_COMB.pep:*
_6/ptodata/2/paa/US08_NEW_COMB.pep:*
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US-09-297-181-4

US-09-297-181-2

US-08-983-035-38

US-09-468-029-58

US-09-468-029-60
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/paa/US087_COMB.pep:
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                                                                                                                  Description
                  Sequence 4, Appli
Sequence 2, Appli
Sequence 38, Appl
Sequence 58, Appl
  Sequence
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ALIGNMENTS

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RESULT 1
US-09-297-181-4
US-09-297-181-4
Sequence 4, Application US/09297181
GENERAL INFORMATION:
APPLICANT: Bracco, Laurent
APPLICANT: Bracco, Laurent
APPLICANT: Debussche, Laurent
TITLE OF INVENTION: ANTI-P53 SINGLE-CHAIN ANTIBODY FRAGMENTS AND THEIR USES
FILE REFERENCE: S196030-US
CURRENT APPLICATION NUMBER: US/09/297,181
CURRENT FILING DATE: 1999-04-26
EARLIER APPLICATION NUMBER: PCT/FR97/01921
EARLIER FILING DATE: 1997-10-27
EARLIER FILING DATE: 1997-10-27
EARLIER FILING DATE: 1996-10-29
NUMBER OF SEO. ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
SEO ID NO 4
LENGTH: 249
TYPE: PRT
ORGANISM: Mus musculus
US-09-297-181-4

Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-108;
Matches 249; Conservative 0; Mismatches 0; Indels 0; Gaps 0:
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RESULT 3
US-08-983-035-38
; Sectione 38. Application US/08983035
; GENERAL INFORMATION:
CONSEILLER, Emmanuel
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GENERAL INFORMATION:
APPLICANT: Bracco, Laurent
APPLICANT: Debussche, Laurent
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Best Local Similarity
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LENGTH: 243
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CURRENT FILING DATE: 1999-04-26
EARLIER APPLICATION NUMBER: PCT/FR97/01921
EARLIER FILING DATE: 1997-10-27
EARLIER FILING TON NUMBER: FR96/13176
EARLIER APPLICATION NUMBER: FR96/13176
EARLIER FILING DATE: 1996-10-29
NUMBER OF SEO ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ANTI-P53 SINGLE-CHAIN ANTIBODY FRAGMENTS AND THEIR USES FILE REFERENCE: ST96030-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.0
APPLICANT:
                                                                                                                                                                     240 FGGGTKLEIK 249
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                                                US-09-468-029-58; Sequence 58, Application US/09468029; GENERAL INFORMATION:
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Best Local Similarity
Matches 186; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FR 95/08729
FILING DATE: 19-JUL-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO FR96/0111
FILING DATE: 17-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: SAVILZKY ESG., MATTIN F.
REGISTRATION NUMBER: 29,699
REGISTRATION NUMBER: 29,699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (610) 454-380 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPUTER:
OPERATING SYSTEM: PC-DOS_MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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              TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                     236 FGAGTKLELK 245
                                                                                                                                                                                                                                                  176
                                                                                                                                                                                                                                                                                                                              117
                                                                                                                                                                                                           240 FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                        181 QKPGQSPKVL1YWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
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                                                                                                                                                                                                                                                                                                                                                                                                                                 61 ADNEQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                  QRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLT
                                                                                                                                                                                                                                                                                                                              SSGGGGSGGGGGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDS-DGKTYLNWLL
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r: 500 Arcola Road, Mailstop 3C43
Collegeville
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P53 PROTEIN VARIANTS AND THERAPEUTICAL
            B7-Binding molecules for treating immune
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy TITLE OF INVENTION:

Floppy disk

diseases

COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

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US-09-468-029-60
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            FILING DATE:
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
LENGTH: 556 amino acids
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                 TITLE OF INVENTION: B7-Bindir TITLE OF INVENTION: diseases NUMBER OF SEQUENCES: 80
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                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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amino acid
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73.5%;
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Pred. No. 6.
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                                                                                                                                                                                                                                                                                                                                SOFTWARE: WordPerfect
SEQ ID NO 5
LENGTH: 242
TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                     Query Match
Best Local Similarity
Matches 175; Conserv
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Best Local Similarity
Matches 181; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Barbera-Guillem, Emilio
APPLICANT: Nelson, M. Bud
TITLE OF INVENTION: Vaccine formulations and methods for immunizing a
TITLE OF INVENTION: individual against shed antigen-specific B cells
                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US OF PRIOR FILING DATE: 1999-06-16
NUMBER OF SEQ ID NOS: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/594,985A CURRENT FILING DATE: 2000-09-12
                                                                                                                                                                                                                                                                                                    FEATURE:
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MOLECULE TYPE: pro
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                    LOCATION:
                121
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                                                                            61 APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                      3 KLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEYAP 62
SSGGGGSGGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ
                                              NEKFKGKATLTADKSSSTAYMQLNSLTSEDSAVYFCKRSYY-----GHWGQGTTVTV 112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KFEGKATLTVDKSSSTAYMQLESLTSEDSAVYYCARFAYYGDYYYIMDYWGQGTTVTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLPTFGG
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                                                                                                                                                                                         70.2%; Score 923.5; 70.0%; Pred. No. 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                 us 60/139,521
                                                                                                                                                                        29;
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                                                                                                                                                                                                      DB 19;
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APPLICANT: Witte, Larry

APPLICANT: Witte, Larry

FILE OF INVENTION: Antibodies Specific to KDR and Uses Thereof

FILE REFERENCE: Sequence Listings 1-34 for 381-71

FURRENT APPLICATION NUMBER: US/09/559,019

CURRENT FILING DATE: 2000-04-25

NUMBER OF SEQ ID NOS: 34

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 28

LENGTH: 238

TYPE: PRT

ORGANISM: Mouse
US-09-559-019-28
            Sequence 27, Application US/09550019

GENERAL INFORMATION:
APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
ITILE OF INVENTION: Antibodies Specific to KDR and Uses Thereof
FILE REFERENCE: Sequence Listings 1-34 for 381-71
CURRENT APPLICATION NUMBER: US/09/559,019
CURRENT FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 27
LENGTH: 240
TYPE: PRT
CRGANISM: Mouse
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US-09-559-019-28
; Sequence 28, Application US/09559019
; GENERAL INFORMATION:
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US-09-559-019-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 173 QKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQYYSYPLT 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    233 FGAGTKLVLK 242
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APNFQGKATVTADTSSNTAYLHLSSLTSEDTTYYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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                                                                                                                                                                                                                                                                                                                                                                                 FGSGTKLEIK 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                            QKPGTSPKLWIYSTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPFT 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSGGGGSGGGGGGGDIELTQSPAIMSASPGEKVTITCSASSSV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QVKLQQSGAELVGSGASVKLSCTTSGFNIKDFYMHWVKQRPEQGLEWIGWIDPENGDSDY 60
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US-09-594-985A-4
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SEQ ID NO 4
LENGTH: 262
TYPE: PRT
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Best Local Similar; ty 70.4
Matches 176; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/594,985A CURRENT FILING DATE: 2000-09-12 PRIOR APPLICATION NUMBER: US 60/139,521
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TITLE OF INVENTION:
FILE REFERENCE: B-2
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APPLICANT: Nelson, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial sequence FEATURE: NAME/KEY:
                                       193
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240 FGGGTKLEIK 249
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                                                                                                                                                                                                           61 APNEQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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                                     QKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQYYSYPLT
                                                                          QKPGQSPKVL1YWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T 239
                                                                                                             SSGGGSGGGGGGGDIVMSQSPSSLPVSVGEKVTLSCKSSQSLLYSGNQKNYLAWYQ
                                                                                                                                                    SSGGGGSGGGGGGGDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                         NERFKGKATLTADKSSSTAYVQLNSLTSEDSAVYFCTRSL----NMAYWGQGTSVTV 132
                                                                                                                                                                                                                                                                   QVQLQQSDAELVKPGASVKISCKASGYTFTDHAIHWVKQNPEQGLEWIGYFSPGNDDFKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FGSGTKLEIK 238
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                                                                                                                                                                                                                                                                                                                                          31;
                                                                                                                                                                                                                                                                                                                                          Score 906; DB 19;
Pred. No. 2.5e-72;
1; Mismatches 39
                                                                                                                                                                                                                                                                                                                                                                               DB 19;
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253 FGAGTKLVLK 262

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; LOCATION:
; OTHER INFORMATION: synthesized
US-09-594-985a-6
                                                                                                                                                                                                                                                                                                                                    US-09-594-985A-7
SOFTWARE: WO
SEQ ID NO 7
LENGTH: 272
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LENGTH: 271
TYPE: PRT
ORGANISM: Artificial sequence
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APPLICANT: Nelson, M. Bud
TITLE OF INVENTION: Vaccir
TITLE OF INVENTION: indivi
                                                                                        CURRENT APPLICATION NUMBER: US/09/594,985A CURRENT FILING DATE: 2000-09-12 PRIOR APPLICATION NUMBER: US 60/139,521 PRIOR FILING DATE: 1999-06-16
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CURRENT FILING DATE: 2000-09-12
PRIOR APPLICATION NUMBER: US 60/139,521
PRIOR FILING DATE: 1999-06-16
                                                                                                                                                                                                  APPLICANT: Barbera-Guillem, Emilio APPLICANT: Nelson, M. Bud TITLE OF INVENTION: Vaccine formula TITLE OF INVENTION: individual aga.
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SOFTWARE: WordPerfec
                                                                NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          135 SSGKTTPPVGGGGGGGGGGGGGGGDIVMSQSPSSLPVSVGEKVTLSCKSSQSLLYSGNQK 194
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      234 SYNLP-TFGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    195 NYLAWYQQKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQ 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                174 NYLAWYQQKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQ 233
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les 172; Conserv
                                                                                                                                                                                REFERENCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SS------GGGGSGGGGGGGGGDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRK 173
                                                WordPertect
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US-09-468-029-68; Sequence 68, Applicat; GENERAL INFORMATION:
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                                                                                                    Query Match
Best Local Similarity
Matches 173; Conserv
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Best Local Similarity
Matches 172; Conserv
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LOCATION:
                                                                                                                                                                                                                                                                                                                                      FILING DATE: *
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                           MOLECULE NO TENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SERVELLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSIEM: PC-DOS/MS-DOS
OPERATING SYSIEM: PC-DOS/MS-DOS
OPERATING SYSIEM: PC-DOS/MS-DOS
OPERATING SYSIEM: PC-DOS/MS-DOS
OPERATING SYSIEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial sequence
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                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  173
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61 APNFQGKATYTADISSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                   23
                                                                                                                                                                                                                                                                   STRANDEDNESS:
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 APNFQGKATYTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                  1 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
                                                                                                                                                                                                                                                                                                        TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                     LENGTH:
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                                   QVQLQQSGPELEKPGASVKISCKASGYSFTGHNMNWVKQSNGKSLEWIGIIDPYYGGTSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QYYSYPLTFGAGTKLVLK 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KNYLAWYQQKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KNYLAWYQQKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCK 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SSAKTTPPSVGGGGGGGGGGGGGGDIVMSQSPSSLPVSVGEKVTLSCKSSQSLLYSGNQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QVQLQQSDAELVKPGASVKISCKASGYTFTDHAIHWAKQKPEQGLEWIGYISPGNDDIKY 82
                                                                                                                                                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/09468029
                                                                                                                                                                                                                                                                                                                         268 amino acids
                                                                                                     Conservative
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                                                                                                                                                                                                                                                    protein
                                                                                                                        66.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B7-Binding molecules for treating immune diseases.
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                                                                                                       22;
                                                                                                                          Score 880; DB 18; Pred. No. 5.1e-70;
                                                                                                         Mismatches
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                                                                                                                                           Length 268;
                                                                                                         Indels
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                                                                                                       Gaps
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                                     82
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                                                                                                                           Query Match
Best Local Similarity
Matches 169; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16, Application US/09364088 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 0: FILING DATE: 07-JUNE-1996 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 0: FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/364,088
                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: MX.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: YASHWART M. Deo, et al.
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
NUMBER OF SEQUENCES: 16
              338
                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION: NAME: Remillard, Jane E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                      278
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                             62 PNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDG-----YALDYWGQGT 116
                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                   GGGTKLEIK 249
                PKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCN-----EGTPTGPYYFDYWGQGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGGTKLEIK 261
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28 State Street, 24th Floor
                                                                                                                                                                                                                                                                                                : (617)227-7400
(617)742-7414
                                                                                                                           Conservative
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                                                                                                                                          66.7%;
                                                                                                                                                                                                                                                                                    16:
                                                                                                                        29;
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                                                                                                                          Score 877; DB 17;
Pred. No. 2.4e-69;
9; Mismatches 36;
                                                                                                                                                       Length 553;
                                                                                                                          Indels 20;
                                                                                                                           Gaps
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APPLICANT:
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Chezian Somasundaram

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RESULT 15
US-09-188-082-16
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US-09-523-279-16
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SEQ ID NO 16
LENGTH: 553
TYPE: PRT
                                                    Sequence 16, Application US/09188082 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/18
PRIOR FILING DATE: 1998-11-06
PRIOR APPLICATION NUMBER: 08/66
PRIOR FILING DATE: 1996-07-07
PRIOR APPLICATION NUMBER: 08/48
PRIOR FILING DATE: 1995-07-07
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1999-
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CURRENT FILING DATE: 2000-03-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                       237 LP-TFGGGTKLEIK 249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     278 IKLQQSGAELVRSGTSVKLSCTASGFNIKDSYMHWLRQGPEQGLEWIGWIDPENGDTEYA
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                                                                                                                                                                                                                                                                                                                                         TVTVSSGGGGSGGGGGGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYL 176
                                                                                                                                                                  YPLTFGAGTKLELK 517
                                                                                                                                                                                                                                             HWFQQKPGTSPKLWIYSTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSS
                                                                                                                                                                                                                                                                              AWYQQKPGQSFKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYN 236
                                                                                                                                                                                                                                                                                                                                                                                              PKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCN-----EGTPTGPYYFDYWGQGT
                                                                                                                                                                                                                                                                                                                                                                                                                                PNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDG-----YALDYWGQGT 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            169;
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Joel Goldstein
Robert Graziano
                                      Yashwant M. Deo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 877; DB 19;
Pred. No. 2.4e-69;
9; Mismatches 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 553;
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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-09-188-082-16
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TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino
TYPE: amino
                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 169; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: MX TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/188, FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/661,052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                    237 LP-TFGGGTKLEIK 249
                                                                                         177 AWYQQKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYN 236
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Bracco,L., Schweighoffer,F. and Tocque,B.
CONDITIONAL EXPRESSION SYSTEM

Patent: WO 9630512-A 4 03-OCT-1996;
RHONE POULENC RORER SA (FR)
Other publication AU 5402096 961016
Other publication FR 2733348 961004.
Location/Qualifiers
               ctgcacctcagcagcctgacatctgaggacaccaaccgtctattactgtaatgcagtcatc
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                                                                             Caron de Fromentel,C., Gruel,N., Venot,C., Debussche,L., Conseiller,E., Dureuil,C., Teillaud,J.L., Tocque,B. and Bracco,I Restoration of transcriptional activity of p53 mutants in human tumour cells by intracellular expression of anti-p53 single chairs are transcriptional activity.
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                                                                                                             60.8%; nilarity 78.7%; Conservative
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132. 24
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/trans1_table=11
/product="anti-p53 SCFV antibody (scFv421)"
/product="anti-p53 SCFV antibody (scFv421)"
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LDYMGQGTTYTVSSGGGGSGGGGSGVLMTQTPLITLSVTIGQPASISCKSSQSL
LDSDGKTYLNWLLQRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAED
LGVYYCWQGTHSPLTFGAGTKLEIK"
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/db_xref="taxon:10090"
168 c 205 q 180
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/db_xref="taxon:10090"
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/db_xref="taxon:32630"
/cell_line="PAb421"
/cell_type="hybridoma"
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/db_xref="taxon:10090"
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                                                                       CTGCAGCTCAGCCTGGCATCTGAGGACACTGCCGTCTATTATTGTAAT------
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Conselller_E. and Bracco, L.
P53 PROTEIN VARIANTS AND THERAPEUTICAL
Patent: WO 9704092-A 36 06-FEB-1997;
RHONE POULENC RORER SA (FR)
Other publication FR 2736915 970124.
Location/Qualifiers
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/db_xref="taxon:32644"
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Other
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Conseiller, E. and Bracco, L.

P53 PROTEIN VARIANTS AND THERAPEUTICAL

P54 PROTEIN VO 9704092-A 31 06-FEB-1997;
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publication FR 2736915 970124
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GGLEWIGWIDENGDTEYAPKFOGKATWIADTSSNIA LQLSSLASEDTAVYYCNFYG
DALDYWGOGTTYTVYSSGGGGSGGGGSGGGSDYLMTOTPLTLSVTIGQDASISCKSSO
SLLDSDGKTYLNWLLQRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEA
EDLGVYYCGWGCHSPLTFGAGTKLELKRAALLQTRAPAPAPTGAAPAPAPSSFWPLSSS
VPSQKTYGGSVGFRLGFLHSGTAKSVTCTYSPALNKWECQLAKTCPVQLWVDSTPPPG
TRVRAMAIYKQSQHMTEVVRRCPHHERCSDSDGLAPPQHLIRVEGNLRVEYLDDRNTF
                                                                                                                                                        ALKEKLKALEEKLKALVGER"
443 c 458 g 341 t
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EVRVCACPGRDRRTEEENLRKKGEPHHELPPGSTKRALPNNTSSSPQPKKKPLDGDLK
                                                                                                                                                                                                                                                                                                                                   /protein_id="CAA03533.1"
/db_xref="GI:3714714"
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TCGCTTCACAGGCAGTGGATCTGGAACAGATTTCACTCTCACCATCAGCAGTGTGCAGGC
                                             AGGGCAGCCTCCTAAACTGTTGATCTACGGGGCATCCACTAGGGAATCTGGGGTCCCTGA
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of binding to an epitope
Patent: WO 9925818-A 27-MAY-1999;
KUFER PETER (DE); RAUM TOBIAS (DE
LOCATION/Qualifiers
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Eukaryota;
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                                      GTCTCCATCCTCCCTGACTGTGACAGCAGGAGAGAGGTCACTATGAGCTGCAAGTCCAG
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Sequence :74
AX003780
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/db_xref="taxon:10095"
174 c 212 g
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Sutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Pred. No. 6.7e-125;
D; Mismatches 147;
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                                   complement(2720. .3580)
/gene="bla"
                                                          1463. .2720
/note="ColE1 origin"
                                                                                                                                       1360.
                                                                                                                                                             /gene="scFv-215"
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/db_xref="taxon:1895"
                                                                                                                                                                                                                                      'note="encodes core streptavidin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="insert part 1"
/organism="Mus musculus"
/db_xref="taxon:10090"
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/gene="CDR-H1"
                                                                                                'note="encodes 5x histidine tags for IMAC purification"
'418. .1462
                                                                                                                        /gene="scFv-215"
                                                                                                                                                                                                                                                     /gene="scFv-215"
                                                                                                                                                                                                                                                                           /gene="scFv-215"
/note="encodes linker
                                                                                                                                                                                                                                                                                                                   'gene="CDR-L3"
                                                                                                                                                                                                                                                                                                                                                                                gene="CDR-L2"
                                                                                                                                                                                                                                                                                                                                                                                                                      /gene="CDR-L2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /gene="CDR-L1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product≖"immunoglobulin light chain, kappa variable
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/transl_table=11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /gene="scFv-215"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ′gene≃"CDR-H3"
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                                                                                                                                                'note="encodes epitope
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TTCGGTGCTGGCACCAAGCTGGAGCTGAAA
                                               GTGCAGGCTGAAGACCTGGCAGTTTATTACTGTCAGAATGATCATAGTTATCCGTTAACG
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Best Local Similarity Matches 593; Conserv
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gtccctgatcgcttcacaggcagtggatctgggacagatttcactctcaccatcagcagt 660
                                                                                                                                 cagaaaccagggcagtctcctaaagtgctgatctactgggcatccactagggaatctgga
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/gene="bla"
.3712. .4284
./note="intergenic region"
.a 1104 c 1126 g 1007 t
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YSPVTEKHLTGELLTAFLHNMSTTLRKLLTGELLTLASRQQLIDMMEADKVAGPL
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EIGASLIKHW"
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D; Mismatches 133;
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597; Conserv
                                                                                                                                                                                                    Artificial sequence; Y14583
                                                                                                                                  artificial sequence; fusion protein; histidine tagged fusion protein; immunoglobulin heavy chain; immunoglobulin light chain; immunoglobulin antibody; streptavidin; variable region.
   Bifunctional and multimeric complexes single chain antibodies (scfv)
                                                                    Plasmid synthetic construct artificial sequence.
1 (bases 1 to 4354)
                                                                                                                      lactamase; recombinant antibody;
synthetic construct.
                                                                                                                                                                                        Y14583.1 GI:2462912
                                       Little, M.
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                                                     ., Breitling, F., Kontermann, R.,
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Submitted (13-AUG-1997) S. Duebel, Molecular Genetics, Uni
of Heidelberg, Im Neuenheimer Feld 230, 69120 Heidelberg,
Related sequences: Y14584, Y14585, X03591 and Y12778.
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and Dubel,S.
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                                                /Codon_start=1
/transl_table=11
/product="fimmunoglobulin heavy chain, variable domain"
/protein_id="CAA74918.1"
/db_xref="GI:4376152"
/translation="0701Q0SGAELVRPGVSVKISCKGSGYKFTDYATHWVKQSHAKS
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AZWGQGTTVTVS"
                                                                                                                                                                                                                                                                                                                                                                                                                    /product-"scFv-215::core-streptavidin fusion protein"
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/db_xref-"GI:2462913"
/tanslation-"MKYLLPTAAGLLLLAAQPAMAQVOLQQSGAELVRPGVSVKISC
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MELPRLTSDDSAIYYCALLRPFAYWGQGTTVTVSSGGGGGGGGGGGGGGDIELTQSP
SSLSVSAGEKVTMSCKSSQSLLNSGNQNNDLAWYQQRBGQRFKLLIYGASTREGGVPD
'274. .288
/gene="CDR-H1"
/product="Vh hypervariable domain
                                                                                                                                                                                                                                                                                                         /gene-
184. .
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GITGTWYNQLGSTFIVTAGADGALTGTYESAVGNAESRYVLTGRYDSAPATDGSGTAL
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118. .
                                                                                                                                                                                      <184. .>525
/gene="Vh"
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/db_xref="taxon:10090"
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/organism="Mus muscul
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/db_xref="taxon:1895"
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/plasmid="psTE-215"
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                                                                                     GTCAAAGTGTTTTATACAGTTCAAATCAGAAGAACTACTTGGCCTGGTACCAACAGAAAC, 541
                                                                                                                                                            agtotocatottocotygotytytoagcaggagagaggtogotatgagotycaaatoca.487
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Bosman, A. and Lorre, K.
B7-BINDING MOLECULES FOR TREATING IMMUNE DISEASES
Patent: WO 9858965-A 30-DEC-1998;
BOSMAN ALFONS (BE); LORRE KATRIEN (BE)
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Single chain antibody
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           artificial sequence.
1 (bases 1 to 864)
Nardone, F., Spano, F. and Crisanti, A.
Direct Submission
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811. .note="encodes C-myc peptide"
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427. 471
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antibody against Plasmodium berghei Pbs21 antigen"
/codon_start=1
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/db_xref="taxon:32630"
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                              TACACGTTCGGAGGGGGCACCAAGCTGGAAATCAAA
      ccgacgttcggcgggggcaccaagctggaaatcaaa
                                                               aaatccagtcagagtctgttcaaccagtagaacccgaaagaattacttggcttggtatcag
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/evidence=experimental
397. .750
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/note="Myc-tag marker
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Pred. No. 4.8e-140;
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Emery, S.C. and Blakey, D.C.
CHEMICAL COMPOUNDS
PATENT: WO 9851787-A 19-NOV-1998;
EMERY STEPHEN CHARLES (GB); ZENECA
LOCATION/Qualifiers
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CNRS EP118, Faculte des
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                                /evidence=experimental
/product="single chain antibody scFv"
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QCLEMIGRINPANGITTYDFKFQGKATITADTSSNTAYLQLSSLTSEDTAVYFCTRVP
YFDYWGQGTTYTVTVSSGGGGSGGGGGGGSDIELTQSPSSLAVTAGEKYTMRCKSSQS
LLWSVNQKNYLSWYQQKEGQPFKLIIYGASIREPWPPDRFTGSGSGTDFTLTINNVHV
ADLAVYYCOHNHGSFLPYTFGGGTKLEIKRAAABQKLISEEDLN"
                                                                                                                                                                                                /function="specifically oxidase subunits VIac"
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/note="heavy cha
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                                      teeteaggtggaggeggtteaggeggaggtggetetggeggtggeggateggaeattgag
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        gtccctgatcgcttcacaggcagtggatctgggacagatttcactctcaccatcagcagt 660
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Patent: WO 9851787-A 19-NOV-1998;
EMERY STEPHEN CHARTER (77)
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/db_xref="taxon:32644"
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Emery,S.C. and Blakey,D.C.
CHEMICAL COMPOUNDS
PATENT: WO 9851787-A 19-NOV-1998;
EMERY STEPHEN CHARLES (GB); ZENECA
LOCATION/Qualifiers
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Similarity 84.3%;
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/db_xrer="taxon:32630" /cell_type="hybridoma" /focus /focus /focus /focus /focus /focus /focus /focus /focus /focus /focus /note="synthetic" /codon_start=1 /product="anti-53 ScFv antibody (scFv1lD3)" /protein_id="CaA10386.1" /protein_id="CA103868" /translation="QYKLQESGAEIVRSGASVNLSCTASGENIKDYYMHWVKQRPEEG LEWIGYIDPESGETENAPNFOGKATYTADTSSNTAYLHLSSLTSEDTTYYYCNAVIYY EYDGYALDYWQGTTYTVSSGGGGGGGGGGGGGTLETGOSPSSLAVSAGEKVAMSC KSSQSLFNSRTRKNYLAWYQQKFGGGFKVLIYWASTRESGVPDRFTGSGSGTDFTLTI SSVQAEDLAVYYCKQSYNLPTFGGGTKLEIK" source 1. 747	0.0000000000000000000000000000000000000	ACO131533 747 bp m Synthetic construct for a containing linker sequenc AJ131533 AJ131533 GI:4033867 antibody; immunoglobulin synthetic construct. synthetic construct artificial sequence. 1 (bases 1 to 747) Caron de Fromentel.C., Gr Conseiller, E., Dureuil.C. Restoration of transcript tumour cells by intracell Fv fragments	397.4 53.2 729 12 AXO18532 AXO185 397.4 53.2 729 12 AXO18608 AXO186 397.4 53.2 1467 12 AXO12780 AXO27 397.4 53.2 1807 12 AXO02779 AXO027 395.8 53.0 729 12 AXO02778 AXO027 395.2 52.9 732 12 XXU44796 U44796	22 428.6 57.4 843 81 A83223 23 428.6 57.4 1998 81 A83230 24 427.2 57.2 729 11 AF002242 25 426 57.0 747 12 AF226875 26 424 56.8 924 81 A62074 27 424 56.8 924 81 A62074 28 424 56.8 924 81 AR088013 30 411.2 55.0 3991 12 ASY12778 31 409.6 54.8 747 11 MMYHVKFG 32 407 54.5 789 13 AX001527 34 401.6 53.6 3970 12 AF162709 35 400.6 53.6 3970 12 ASY14585 36 400.6 53.6 3970 12 ASY14586 37 400.6 53.6 4864 12 ASY14586 38 400.6 53.6 4864 12 ASY14586 39 10 5526 11 MMSCFVP25 31 400.6 53.6 4864 12 ASY14586 31 400.6 53.6 4864 12 ASY14586 32 400.6 53.6 4864 12 ASY14586 33 402.8 53.9 4363 12 SCFV18290 34 401.6 53.6 4864 12 ASY14586 35 400.6 53.6 4864 12 ASY14586 36 400.6 53.6 4864 12 ASY14586 37 400.6 53.6 4864 12 ASY14586 38 400.6 53.6 4864 12 ASY14586
Qy 541 cagaaaccagggcagtctcctaaagtgctgatctactgggcatccactagggaatctgga 600	Qy 361 tcctcaggtgjaggcggttcaggcggaggtggctggcggttgcggatcggacattgag 420	121 cctgaagagggcctggagtggattggatatattgatcctgagagtggtgaaactgaatat 1	Query Match 100.0%; Score 747; DB 12; Length 747; Best Local Similarity 100.0%; Pred. No. 2.7e-208; Matches 747; Conservative 0; Mismatches 0; Indels 0; Gaps Qy 1 caggicaggagicaggagicaggacitigili	/db_xref="taxon:10000" V_region

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Listing first 45 summaries
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SUMMARIES

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435.4	438.4	442.4	446	447.8	448.4	452.8	454.4	457.6	457.6	462.8	466	469.2	471	501.6	510.2	514.4	548.6	548.6	548.6	747	Score
58.3	58.7	59.2	59.7	59.9	60.0	60.5	60.3	61.3	61.3	62.0	62.1	62.8	63.1	67.1	68.3	68.J	73.1	73.4	73.4	100.0	Query
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AX001519 Sequence	AR054409 Sequence	A45006 Sequence 1	A68604 Sequence 4	AF169027 Synthetic	AX003782 Sequence	A57272 Sequence 4	AJ131532 Synthetic	A59381 Sequence 31	A59386 Sequence 36	AX003780 Sequence	AX003772 Sequence	Y14583 Artificial	U31739 Single chai	AX001511 Sequence	AX001509 Sequence	AJ131195 Mus muscu	A83217 Sequence 37	A83211 Sequence 31	A83227 Sequence 47	AJ131533 Synthetic	Description

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Sequence 1, Appli Sequence 11, Appl Sequence 11, Appl Sequence 14, Appl Sequence 7, Appli Sequence 7, Appli Sequence 5, Appli Sequence 5, Appli Sequence 148, App Sequence 148, App Sequence 148, App Sequence 148, App Sequence 148, App Sequence 148, App Sequence 34, Appli Sequence 34, Appli Sequence 34, Appli Sequence 34, Appli

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2: /cgn2_6/ptodata/2,
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212.818 Million cell updates/sec
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TOPOLOGY: :linear
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                                                                                                                         Matches
                                                                                                                                          Query Match
Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 270 amino acid
TYPE: amino acid
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CITY: Arlington
STATE: Virginia
TANTRY: USA
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 25,327
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703.816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/652,507
FILING DATE: 02-U1-1996
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Arthur R. Crawford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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COUNTRY:
ZIP: 222
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61 APNFQGKATVIADTSSNTAYLHLSSLTSEDTTYYYCNAVIYYEYDG-----YALDYWGQG
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US-08-465-4738-5
US-08-48-1138-148
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US-08-48-16-360-148
US-08-48-360-148
US-08-131-398-148
US-08-131-398-34
US-08-331-398-34
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Result No.

883 877

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Score 883; DB Pred. No. 7.6e 28; Mismatches

DB 2; I 1.6e-67; nes 36;

Length 270; Indels

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Gaps 60

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Post-processing:

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Best Local Similarity
Matches 170; Conserv
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                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: August 14
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/1
FILING DATE: June 16, 1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: WILSON, Steve
TITLE OF INVENTION: A BIFUNCTIONAL OR BIVALENT
TITLE OF INVENTION: ANTIBODY FRAGMENT ANALOGUE
                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANDRESSEE: PILLSBURY, MADISON & SUTRO LLP STREET: 1100 New York Avenue, N.W. STREET: 9th Floor, East Tower CITY: WASHINGTON, D.C. STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS DOS Tes
SOFTWARE: MS WORD, WINDOWS 95, Ver. 4.0
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                       SSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                        QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LAWYQQKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSY 235
SSGGGGSGGGGGGGGSDIELTQSPFSLTVTAGEKVTMNCKSGQSLLNSVNQRNYLTWYQ
                                                                      SDNVKGRFTISRDNGKNTLYLQMSSLKSEDTAVYYCARHGYYG-KGY-FDYWGQGTTVTV 140
                                                                                                    APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                                                     QVQLQESGGDLVKPGGSLTLSCATSGFTFSSYAFSWVRQTSDKSLEWVATISSTDTYTYY 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NLP-TFGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MHWFQQKPGTSPKLWIYSTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRS 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTVTVSSGGGGSGGGSGGGSENVLTQSPAIMSASPGEKVTITCSASSSV-----SY 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTVTVSSGGGGSGGGGGGGGGGDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNY 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCN-----EGTPTGPYYFDYWGQG 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20005-3918
                                                                                                                                                                                                                                                                                                                                   H: 282 amino acids amino acid
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                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VAN DER LOGT, Cornelis Paul Erik
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                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                    protein
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UMBER: PCT/EP/96/03605

August 14, 1996

EQ ID NO: 10:
                                                                                                                                                                                                                     66.7%;
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                                                                                                                                                                                                    20; Mismatches
                                                                                                                                                                                                    Score 877.5; DB 2;
Pred. No. 2.3e-66;
0; Mismatches 57;
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US-08-661-052-16
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                                                                                                        338
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; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-661-052-16
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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LENGTH: 553 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MX
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/661,052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES NUMBER OF SEQUENCES: 16
                    177 AWYQQKPGQSFKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYN
                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 02109-1875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
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                                                                                                                                                                                                                                    TVTVSSGGGGGGGGGGGSENVLTQSPAIMSASPGEKVTITCSASSSV-----SYM 443
                                                                                TVTVSSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYL 176
                                                                                                                                    PKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCN-----:EGTPTGPYYFDYWGQGT 390
                                                                                                                                                                          PNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDG-----YALDYWGQGT 116
                                                                                                                                                                                                              IKLQQSGAELVRSGTSVKLSCTASGFNIKDSYMHWLRQGPEQGLEWIGWIDPENGDTEYA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FGGGTKLEIK 270
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6, Application US/08661052
5837243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   553 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chezian Somasundaram
                                                                                                                                                                                                                                                                                             Conservative
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Robert Graziano
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                                                                                                                                                                                                                                                                                                             66.7%; Score 877; DB 2; Length 553; 66.5%; Pred. No. 5.8e-66;
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Suite 510
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                      236
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444 HWFQQKPGTSPKLWIYSTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSS 503

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                                                                                                                           Matches 164;
                                                                                                                                          Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-7AN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FR93/00085
                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                  LENGTH: 249 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US C. APPLICATION NUMBER: US C. 28-JUL-1994

""MRER: FR 92/01064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: UZIP: 19426
64 NGKFKGKATLTADRSSSTAYMQLSSLTSVGSAVYFCAKENNR--FDERGYYAMDYWGQGT 121
                              61 APNFQGKATYTADTSSNTAYLHLSSLTSEDTTVYYC----NAVIYYEYDGYALDYWGQGT 116
                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                       NAME: Smith Ph.D., Julie K. REGISTRATION NUMBER: P-38,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                             1 QVKLQESGAELVRSGASVNLSCTASGENIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60
                                                             QVQLEQSGPELVKPGASVKISCKASGYAFSRSWMNWVKQRPGQGLEWIGRIYPGDGDTKY 63
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5876969
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Rhone-Poulenc Rorer Inc. 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                   249 amino acids
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Yeh, Patrice
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Guitton, Jean-Dominique
                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                         (610)
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                                                                                                                                                                                                                                                                                                                                        (610)
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64.6%;
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ER: ST92006-US
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                                                                                                                        39;
                                                                                                                                       Score 842.5; DB 2
Pred. No. 1.7e-63;
                                                                                                                           Mismatches
                                                                                                                                                        DB 2;
                                                                                                                           38;
                                                                                                                                                        Length
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US-08-256-790-2
                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                  Matches 165;
                                                                                                                                                                                                                                                                                                                       TELEFAX: 703-243-6410 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/256,790
FILING DATE: :22-JUL-1994
CIASCIPCATION.
                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 9
FILING DATE: '23-JAN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/
FILING DATE: 15-JAN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: PLUECKTHUN, APPLICANT: PACK, PETER
                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.
ZIP: 22201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                236 YPWTFGGGTKLEIK 249
                      59 EYAPNEQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTV 118
                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: MEI
                                                                                                                                                                                                                                                                                                                                                                                                          NAME: HAMLET-KING, DIANA REGISTRATION NUMBER: 33,
 82
                                                                 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                     TOPOLOGY:
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                                                                                                                                                                                                                                                                                     LENGTH:
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                                                                                   1 QVKLQESGAELVRSGASVNLSCTASGENIKDYYMHWVKQRPEEGLEWIGYIDPESGE--T 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TVTVSSGGGGSGGGGGGGSNIQLTQSPNSMSTSVGDRVSITCKASQDVDTS-----V
EYSASVKGRFIVSRDTSQSILYLQMNALRAEDTAIYYC-ARNYYGSTWY-FDVWGAGTTV 139
                                                                 EVKLYESGGGLYQPGGSLRLSCATSGFTFSDFYMEWVRQPPGKRLEWIAASRNKGNKYTT
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                                                                                                                                                                                                                                                                   amino acid
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                                                                                                                                                                                                                                                                                       277 amino acids
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                                                                                                                                  Conservative
                                                                                                                                                                                                                                                     Linear
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                                                                                                                                             63.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MONOMERIC AND DIMERIC ANTIBODY FRAGMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Release #1.0, Version #1.30
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                                                                                                                                                 Score 831.5; DB 2
Pred. No. 1.6e-62;
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/D. SUITE 1400
                                                                                                                                 Mismatches
                                                                                                                                  53;
                                                                                                                                                                  2;
                                                                                                                                  Indels
                                                                                                                                                                Length 277;
                                                                                                                                  5.
                                                                                                                                 Gaps
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                          TELEFAX: 703-243-6410 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: METELECOMMUNICATION INFORMATION: 703-243-6333
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                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: EP 9.
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: PIULATS
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                             APPLICATION NUMBER: EP 94118970.6
FILING DATE: 02-DEC-1994
ATTORNEY_AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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260 LTFGAGTKLELK 271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 17-NON CLASSIFICATION: 530
                                                                                                     TOPOLOGY:
                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                               REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: WO PCT/EP95/00978 FILING DATE: 16-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: ARLINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            YQQKPGQPPKLLIYGASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCQNDHSYP 259
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                                                                                                                                    244 amino acids
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ROSELL, ELISABET
BLASCO, FRANCESC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PIULATS, JAUME
PENTION: ANTI-EGER SINGLE-CHAIN FVS AND ANTI-EGER
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17-NOV-1995
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 62.6%;
63.1%;
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                                                                                                                                                                                                                                                       33,302
 Score
Pred.
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 823.5; DB 2;
No. 6.5e-62;
                  Length 244;
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                                                                                      ; MOLECULE TYPE: US-08-661-052-14
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Patent No. 5837243
GENERAL INFORMATION:
                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/48.

FILING DATE: 07-JUNE-1995

ATTORNEY_AGENT INFORMATION:

NAME: AITOOLd, Beth E.

REGISTRATION NUMBER: MXI-

REFERENCE/DOCKET NUMBER: MXI-

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                           TELEFAX: (617)227-594
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chezian Somasundaram TITLE OF INVENTION: THERAPEUTIC TITLE OF INVENTION: OF ANTI-FC
                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 02109-1875
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                                                                                                                                       amino acid
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                                                                                                                                                          301 amino acids
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                                                                                                                                                                                                           : (617)227-7400
(617)227-5941
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Robert Graziano
                                                                                                                          linear
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                                                                                                    protein
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62.2%; Score 818; DB 2; 62.1%; Pred. No. 2.4e-61; tive 37; Mismatches 55
                                                                                                                                                                                                                                                                                                                                                                                                             US/08/661,052
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                               Length 301
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Matches 154;

Conservative

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US-08-553-497A-26
                    TELEFAX: 703-243-6410 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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itent No. 584***
rear information:
xettle
                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                              RECISTRATION NUMBER: 33,302
REFERENCE/DOCKET NUMBER: MEI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-243-6333
                                                                                                                                                       APPLICATION NUMBER: EP 94
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
                                                                                                                                                                                                                                   APPLICATION NUMBER: EP 9/
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                               FILING DATE: 16-MAR-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 17 CLASSIFICATION:
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                                                                           TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KPGKAPKLLIYWASTRESGVPSRFSGSGSGTDFTFTISSLQPEDIATYYCHQYLSSWTFG
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ROSELL, ELISABET
BLASCO, FRANCESC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
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                                                                                                                                                                                                                                                                                 EP 94104160.0
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US-08-553-497A-24
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                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NYMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24, Appl Patent No. 584409
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Best Local Similarity
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                                                       APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94104160.0
FILING DATE: 17-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN, P.C
STREET: 2200°CLARENDON BLVD. SUITE 1400
                                    FILING DATE: 17-MAR-PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US .
ZIP: 22201
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APPLICATION NUMBER: EP 94118970.6 FILING DATE: ,02-DEC-1994
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GY: linear
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ROSELL, ELISABET
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Pred. No. 4.4e-61;
8; Mismatches 53;
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SEQUENCE CHARACTERISTICS:
LENGTH: 246 amino acids
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tent No.
                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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NAME: HAMLET-KING, DIANA
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                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
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APPLICATION NUMBER: US/08/553,497A FILING DATE: 17-NOV-1995
                                                                                                                                                        COUNTRY:
                                                                                                                                                                                           CITY: ARLINGTON
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                                                                                                                                         22201
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2200 CLARENDON BLVD.
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//ENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
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Pred. No. 6.6e-61
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Best Local S
                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                        Sequence 19,
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                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                               APPLICANT: Filpula, David
TITLE OF INVENTION: Multivalent Antigen-Binding Proteins
NUMBER OF SEQUENCES: 23
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ATTORNEY/AGENT UNFORMATION:
                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          173
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                                                                                                                                                                                                                                                                                                                                                                                                          233 YTFGGGTKLEIK 244
                                                                                                                                                                                                                                                                                                                                                                                                                                       239 -TFGGGTKLEJK 249
                                                                                             CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 703-243-6410
                                                  COUNTRY: U.S.A.
ZIP: 20005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: HAMLET KING, DIANA REGISTRATION NUMBER: 33,302 REFERENCE/DOCKET NUMBER: ME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: EP 9
FILING DATE: 17-MAR-1994
                                                                                     STATE:
                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 NEKIKSKATLTVDKSSSTAYMQLSSLTSEDSAVYYC-ASRDYDYDGRYFDYWGQGTTVTV 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: EP 94118970.6 FILING DATE: 102-DEC-1994
                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QKPGQSPKVLIYMASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-- 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SSGGGGGGGGGGGGGGDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EVQLQQSGAELVKPGASVKLSCKASGYTFTSHWMHWVKQRAGQGLEWIGEFNPSNGRTNY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SSGGGGSGGGGSGGGSDIELTQSPTIMSASPGEKVTMTCSDSSSV-----SYMYWYQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           157;
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                                                                                   D.C.
                                                                                                                                                                                                                                                                                                                        Application US/08392338A
                                                                                                                    E: Sterne, Kessler, Goldstein & Fox P.L.L.C 1100 New York Avenue, NW
                                                                                                                                                                                                                                                                       Whitlow, Marc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                      Wood, James F
Hardman, Karl
                                                                                                                                                                                                                       Bird,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-MAR-1995
                                                                                                                                                                                                                     Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61,5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/989,846
FILING DATE: 20-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/796,936
FILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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                                                                                                                 APPLICANT: Bird, Robert
APPLICANT: Filpula, David
APPLICANT: Follence, Michelle
TITLE OF INVENTION: Methods fc
                                                                                                                                                                                                                                                                                      tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 483 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 0977.0030007
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEFAX: (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                346 FGAGTKLVLK 355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             188 NERFKGKATLTADKSSSTAYVQLNSLTŠEDSAVYFCTRSL-----NMAYWGQGTSVTV 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     181 QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
COUNTRY:
                                                STREET:
                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                                                                                                                                                                                    FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQYYSYPLT 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QVQLQQSDAELVKPGASVKISCKASGYTFTDHAIHWVKQNPEQGLEWIGYFSPGNDDFKY 187
                           Washington
                                                                                                                                                                                                                                                                                    9, Application US/09166750
6025165
                  D.C.
                                              E: Sterne, Kessler, Go
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Hardman, Karl
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                                                                                                                       Methods for Producing Multivalent Antigen-Binding Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61.1%; Score 803.5; DB 2 62.4%; Pred. No. 7.1e-60; tive 32; Mismatches 39
                                                  Kessler, Goldstein & Fox P.L.L.C
ork Avenue, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US/08/392,338A
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US-09-166-093-19
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                                                                                                                                   Patent No. 6027725
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 156;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                             Sequence 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 07//yo, APPLICATION NUMBER: US 007//yo, EILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A.
NAME: Goldstein, Jorge A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (202) 371-25
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 483 amino acid
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TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/392,338
FILING DATE: 22-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/989,846
FILING DATE: 20-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Reli
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                 APPLICANT:
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PRIOR APPLICATION DATA:
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                       APPLICANT:
                                                             APPLICANT:
                                                                               APPLICANT:
                                                                                                             APPLICANT: Whitlow, Marc
                                                                                                                                                                                                                                                                                                                                                    286
                                                                                                                                                                                                                                                                346 FGAGTKLVLK,355
                                                                                                                                                                                                                                                                                                            240 FGGGTKLEIK 249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: ami
PPLICANT: Filpula, David
PPLICANT: Rollence, Michelle
ITLE OF INVENTION: Multivalent Antigen-Binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 128 QVQLQQSDAELVKPGASVKISCKASGYTFTDHAIHWVKQNPEQGLEWIGYFSPGNDDFKY 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM.PC compatible
OPERATING SYSIEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60
                                                                                                                                                                                                                                                                                                                                                    QKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQYYSYPLT
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                                                                                                                                                                           Application US/09166093
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                                                                           Wood, James F
Hardman, Karl
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                                                         Bird, Robert
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Gaps

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285

240

NUMBER OF SEQUENCES:

Bird, Robert

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RESULT 14
US-09-172-019-19
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                                                            Sequence 19, Application US/09172019 Patent No. 6103889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 156;
  GENERAL INFORMATION:
APPLICANT: Whitlo
APPLICANT: Hardma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ANTIGNALIZATION NOVINEER: 29,021
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0977.003000B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/796,936
FILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/989,846
FILING DATE: 20-NOV-1992
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 08/392,338
FILING DATE: 22-FEB-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           128
                                                                                                                                                                                                                                                                                                                                                                                                                                           188
                                                                                                                                                                                                                                                                         286 QKPGQSPKLLIYWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQYYSYPLT
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                         FGAGTKLVLK 355
                                                                                                                                                                                                               FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                          QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
                                                                                                                                                                                                                                                                                                                                                       S-----SDVVMSQSPSSLPVSVGEKVTLSCKSSQSLLYSGNQKNYLAWYQ 285
                                                                                                                                                                                                                                                                                                                                                                                                SSGGGGSGGGGGGGGDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QVQLQQSDAELVKPGASVKISCKASGYTFTDHAIHWVKQNPEQGLEWIGYFSPGNDDFKY 187
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Whitlow, Hardman,
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Best Local S
Matches 156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A.
REGISTRATION HUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0977
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 20-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 25-NOV-1991
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FILING DATE: 22-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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MEDIUM TYPE: 'Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 483 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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CLASSIFICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                           1 QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
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                                                                                                                                                                           SSGGGGSGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSOSLFNSRTRKNYLAWYQ 180
FGAGTKLVLK 355
                                   FGGGTKLEIK 249
                                                                           OKPGQSPKLL1YWASARESGVPDRFTGSGSGTDFTLSISSVKTEDLAVYYCQQYYSYPLT
                                                                                                                                                       S-----SDVVMSQSPSSLPVSVGEKVTLSCKSSQSLLYSGNQKNYLAWYQ
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-09-166-094-19
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Best Local Similarity 62.4
Matches 156; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/392,338
FILING DATE: 22-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/989,846
FILING DATE: 20-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/796,936
FILING DATE: 25-NOV-1991
APPLICATION TORNALTION:
APPLICATION TORNALTION:
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TITLE OF INVENTION: Multivalent Antigen-Binding Proteins
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                 181 QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T 239
                                                                                                                                                         121 SSGGGGSGGGGGGGGDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                                                                            128 QVQLQQSDAELVKPGASVKISCKASGYTFTDHAIHWVKQNPEQGLEWIGYFSPGNDDFKY 187
240 FGGGTKLEIK 249
                                                                                                                                                                                                                    188 NERFKGKATLTADKSSSTAYVQLNSLTSEDSAVYFCTRSL-----NMAYWGQGTSVTV 240
                                                                                                                                                                                                                                             61 APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 29, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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D.C.
RY: U.S.A.
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Bird, Robert
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                                                                                                                                                                                                                                                                                                                                                                                              61.1%; Score 803.5; DB 3; Length 483; 62.4%; Pred. No. 7.1e-60; ative 32; Mismatches 39; Indels 23; Gaps
                                                                                                                             -SDVVMSQSPSSLPVSVGEKVTLSCKSSQSLLYSGNQKNYLAWYQ 285
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Db 346 FGAGTKLVLK 355

Search completed: February 12, 2001, 15:23:46
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Perfect score:
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1: /cgn2_2/gcgdata/geneseq/geneseqn/NA1980.DAT:

2: /cgn2_2/gcgdata/geneseq/geneseqn/NA1981.DAT:

3: /cgn2_2/gcgdata/geneseq/geneseqn/NA1982.DAT:
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Gapop 10.0 , Gapext 1.0
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/cgn2_2/gcgdata/geneseq/geneseqn/NA1986.DAT:*
/cgn2_2/gcgdata/geneseq/geneseqn/NA1987.DAT:*
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Mouse scFV fragmen
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Anti-CD86 monoclon
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Fusion protein pIC
Plasmid pNG4/55.1s
Human p53 protein
               Anti-human protein
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ALIGNMENTS

08-SEP-1998 (first entry)

V36237;

V36237 standard; DNA; 747 BP.

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Single chain antibody; ScFv D3M; mouse; p53 protein; oligomerisation; regulatory domain; p53 mutant; H273; W248; G281; p53-dependent trans-activating activity; restoration; tumour-suppressing activity; tumour cell; treatment;
Restoring p53-dependent trans-activating activity to cell containing mutant p53 - by delivering single-chain antibody specific for the
                                                                                      WPI; 1998-272140/24.
P-PSDB; W60770.
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Best Local Similarity
Matches 630; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conjugate; cell targeting; cytotoxic drug; plasmid; fusion protein; prodrug-converting enzyme; cell surface antigen; treatment; cancer; inflammation; rheumatoid arthritis; antibody; prodrug therapy syste
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 864 BP; 214 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New gene construct expressing conjugate of targeting agent and prodrug-converting enzyme - useful for, e.g. targeted production cytotoxic drug in vivo, especially for treatment of cancer
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                                                                                                                                                                                                                                                                       Sequence
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Chimeric
                                                                                                                                                                                                            B7 binding molecule; costimulatory molecule; B7.1; CD80; B7.2; CD86; T cell activation; inhibitor; graft versus host disease; transplant rejection; allograft rejection; autoimmune disease; allergy; therapy; human; bispecific tetravalent antibody; BiTAI BITABB7-24-1G10H6; ss.
                                                                                                                               CDS
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           has been expressed in Escherichia coli cells. It cross-links, and/or cross-reacts, with the costimulatory molecules B7.1 and B7.2 expressed on the membrane of professional antigen-presenting cells, leading to the inhibition of antigen-specific T cell activation. The invention relates to such B7-binding molecules, methods for their production, and their use for treating or preventing diseases of the immune system, in particular graft rejection, graft versus host disease, allergy and autoimmune diseases (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This DNA sequence encodes the bispecific tetravalent antibody BITABB7-24-IG10H6 (see W90217) comprising the anti-B7.1 murine monoclonal antibody (MAb) B7-24 VL region joined, via a flexible synthetic linker, to the B7-24 VL region, the human IgG3 hinge region, a helix-turn-helix dimerisation domain, the human IgG3 hinge domain, the anti-B7.2 MAb IG10 VH region, another flexible linker, the IG10 VL region and a hexahistidine tag. The BITAB has been expressed in Escherichia coli cells. It cross-links,
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                                                       ctcacccagtctccatcttccctggctgtgtcagcaggagagaggtcgctatgagctgc 480
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Pred. No. 1.3e-123;
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rejection or treat autoimmune diseases such as rheumatoid arthritis, multiple sclerosis, psoriasis and systemic lupus erythematosus. It can also be used to inhibit a response to gene therapy vectors or other therapeutic molecules. A single compound can bind to both CD40 and CD86 (expressed at low level by professional antigen presenting cells), resulting in inhibition of T cell activation and inducing T cell anergy.
                                                                                                                    CD86 monoclonal antibody Fun-1 ScFv construct, from an example of the present invention. The present invention describes a soluble molecule that can bind human CD40 antigen and at least human CD86 antigen, which are both present on the surface of lymphocytes. The soluble molecule can be used to induce T cell tolerance and to treat T-cell mediated diseases, specifically to prevent allograft or xenograft transplant
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                                                                                                                                                                                                                                                                                                                                             Soluble compound that binds both CD40 and CD86 on surface of lymphocytes - is used to induce T cell tolerance and treat T cemediated diseases, also recombinant vectors and host cells that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-CD40; monoclonal antibody; human; lymphocyte; T cell; allograft; xenograft transplant rejection; autoimmune disearheumatoid arthritis; multiple sclerosis; psoriasis; ss.
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                                                                                                                                                                                                                                                                                   Example 1; Page 27; 47pp; English.
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Best Loc
Matches
B7 binding molecule; costimulatory molecule; B7.1; CD80; CD86; T cell activation; inhibitor; graft versus host distransplant rejection; allograft rejection; autoimmune disallergy; therapy; human; bispecific tetravalent antibody
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                                                     tetravalent antibody BiTAb1G10-B7-24H6
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Pred. No. 4.4e-122;
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                                                                                                                                                                                                                                                                                                                                                                                     CC This DNA sequence encodes the bispecific tetravalent antibody CC Bi7AbG10-B7-24H6 (see W90218) comprising the anti-B7.2 murine CC monoclonal antibody (MAb) 1G10 VH region joined, via a flexible CC synthetic linker, to the 1G10 VI region, the human 1G3 hinge CC synthetic linker, to the 1G10 VI region, the human 1G3 hinge CC region, a helix'turn-helix dimerisation domain, the human 1G3 and thinge domain, the anti-B7.1 MAb B7-24 VH region, another flexible CC linker, the B7-24 VI region and a hexahistidine tag. The Bi7Ab CC has been expressed in Escherichia coli cells. It cross-links, CC and/or cross-reacts, with the costimulatory molecules B7.1 and B7.2 CEXPRESSED ON The membrane of professional antigen-presenting cells, CC leading to the inhibition of antigen-specific T cell activation. The invention relates to such B7-binding molecules, methods for CC their production, and their use for treating or preventing diseases CC of the immune system, in particular graft rejection, graft versus CC host disease, allergy and autoimmune diseases (claimed).
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Best Local Similarity
Matches 591; Conserv
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Chimeric - 1
Chimeric - 1
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The invention relates to a method of identifying binding site domains (BSD) that retain the capacity of binding to a predetermined epitope when positioned C-terminal of at least one further domain in a recombinant bior multivalent polypeptide. The method comprises (a) testing a pagel of BSD displayed on the surface of a biological display system as part of a fusion protein for binding to a predetermined epitope, where the fusion protein comprises an additional domain positioned N-terminal of the BSD and an amino acid sequence that mediates anchoring of the fusion protein to the surface of the display system; and (b) identifying a BSD that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-NOV-1998;
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Best Local Sim
Matches 585;
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                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method of identifying binding site domains (CC (BSD)) that retain the capacity of binding to a predetermined epitope when compositioned C-terminal of at least one further domain in a recombinant bicom multivalent polypeptide. The method comprises (a) testing a panel of the surface of a biological display system as part of a comprise of the surface of a biological display system as part of a comprise of a protein comprises an additional domain positioned N-terminal of the BSD and an amino acid sequence that mediates anchoring of the fusion protein to the surface of the display system; and (b) identifying a BSD that thinds to the predetermined epitope. The method is useful to identify bicor multivalent polypeptides that comprise antibody binding sites capable of efficiently binding to the corresponding antigen. The polypeptides or antibodies identified by the method are useful therapeutically and cantibodies identified by the method are useful therapeutically and cantibodies identified by the method are useful therapeutically and cantibodies identified by the method are useful therapeutically and cantibodies identified by the method are useful therapeutically and cantibodies identified by the method are useful therapeutically and cantibodies identified by the method are useful therapeutically and cantibody cantibodies identified by the method are useful therapeutically and proteins can be isolated from combinatorial antibody libraries using the new in vitro method. Sequences x77240-248 represent the contraction of the protein scale protein sector of the protein scale protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Phage display system for identification retaining capacity to bind an epitope
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P-PSDB; Y17964.
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gticaagggcaaagccacactgactgcagacaaatcttcgagcacagcctatatgcagct
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                                                                                                                                                                                                                                                                                                                                                                                                                           183 A;
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Pred. No. 2.2e:
0; Mismatches
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Restoring p53-dependent trans-activating activity to cell mutant p53 - by delivering single-chain antibody specific
                                                                                                                                                                                                                                                                     single chain antibody; regulatory domain; p53
                                    WPI; 1998-272140/24.
P-PSDB; W60769.
                                                                                                                                           27-OCT-1997;
                                                                                                                                                                 07-MAY-1998
                                                                                                                                                                                       WO9818825-A1.
                                                                                                                                                                                                              Mus sp
                                                                                                                                                                                                                                    tumour-suppressing activity;
hyper-proliferation; cancer;
                                                                                                                                                                                                                                                         Single chain antibody; ScFv 421; mouse; p53 protein; regulatory domain; p53 mutant; H273; W248; G281; p53-dependent trans-activating activity; restoration;
                                                                                                                                                                                                                                                                                                        DNA of ScFv 421 which binds to mutant p53 proteins
                                                                                                                                                                                                                                                                                                                                                                               V36236
                                                                                                                   29-OCT-1996;
                                                                                            (RHON ) RHONE-POULENC RORER
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cancer; re-stenosis;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 729 BP; 174 A; 171 C; 207 G;
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nes 594; Conservative
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                                                   gtgcaggctgaagacctggcagtttattactgcaagcaa---tcttataatctaccgacg
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                              gtggaggctgaggatttgggagtttattattgctggcaaggtacacattctccgctcacg
                                                                                                    gtccctgacaggttcactggcagtggatcagggacagatttcacactgaaaatcaacaga
                                                                                                                                                                    cagaggccaggccagtctccaaagcgcctaatctatctggtgtctaaactggactctgga
                                                                                                                                                                                                                                          aagtcaagtcagagcctcttgga---tagtgatggaaagacatatttgaattggttgtta
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Pred. No. 7.3e-111;
0; Mismatches 132; Indels
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61 10

Query Match Best Local (Matches

Local Simbles 592;

Similarity

61.3%; 78.9%;

Score 457.6; DB Pred. No. 5e-110;

DB 18;

Conservative

0;

Mismatches 134;

Indels Length

24;

Gaps

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1 caggtcaagctgcaggagtcaggggcagaacttgtgaggtcagggggcctcagtcaatttg

tcctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaaacagagg 120 caggtgcagc;gcaggagtcaggggcagagcttgtggggtcaggggcctcagtcaagttg

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RESULT 11
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                                       Claimed variants of protein p53 have at least part of the coligomerisation domain deleted and replaced by a leucine zipper domain. The mutants preferably also have at least part of the p53 transactivation domain (amino acids 1-74) deleted and replaced by the transactivating domain (TD) from herpes simplex virus viral protein vP16 (amino acids 411-490) or by a protein domain able to bind selectively to a transactivator, especially a single-chain antibody variable domain (ScFv). The present sequence represents the Ncol-NotI fragment of CDNA coding for the anti-p53 single-chain codon cprecise location is not indicated) and no termination codon. The p53 variants are more active and more stable tumour suppressors and apphtosis-inducing agents than wild-type p53 and are active where the wild-type protein is not, i.e. they are not inactivated by contains (because the leucine zipper domain prevents formation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leucine zipper domain; LZD; oligomerisation domain; mutant; substitution; replacement; transactivation; viral protein VI anti-oncogene; hyperproliferation; cancer; restenosis; ScFv
Sequence 749 BP; 173 A; 179 C;
                                                                                                                                                                                                                                                                                                                                                                New p53 variants e.g. with oligomerisation domain replaced by leucine zipper to useful for treating hyper-proliferative disorders.
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anti-oncogene; hyperproliferation; cancer; restenosis;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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The invention relates to novel chimaeric, bispecific proteins which comprise: (a) a DNA binding domain and (b) a domain which binds a trans-activator (TA), trans-repressor (TR) or their complexes, which are characteristic of a physiological or physiopathological state. The novel chimaeric, bispecific proteins allow expression of a therapeutic protein (e.g. diphtheria or Pseudomonas toxins, thymidine kinase, single chain antibodies) to be regulated in response to particular conditions. The Ta- or TR-binding domain may be a antibody such as a single chain
                                                                                                                                                                                                                                                                                                                         Conditional gene expression system triggered by e.g. infection or hyper-proliferation - comprises novel bi-specific proteins having DNA-binding domain and second domain specific for trans-activator of the property of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific for trans-activator of the second domain specific fo
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                                                                                                                                                                                                                                       Query Match
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Matches 574
                                                                                                                                                                                                                                                                                                                 Sequence 744 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Phage display system for identification retaining capacity to bind an epitope
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gttcaagggcaaagccacactgactgcagacaagtcctcgtacacagcctatatgcagct
                                                         tggacttgaatgggttggagatattttccctggaagtggtaatgctcactacaatgagaa
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                                                                                                                                                                                                                                                                                                                               V09256;
26-JUN-1997;
                    31-DEC-1997
                                          W09749726-A1
                                                                                                                                                                                                                                         protein synthesis; conjugate;
fusion protein; chimeric.
                                                                                                                                                                                                                                                                                  Nucleotide sequence encoding the Mgr6-clavin immunotoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1314 BP; 338 A; 360 C; 359 G; 257 T; 0 other;
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1: /cgn2_6/ptodata/2/f

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/cgn2_6/ptodata/2/pna/US089_COMB.seq:*
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6/ptodata/2/pna/US095C_COMB.seq:*
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/ptodata/2/pna/US6010_COMB.seq
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1509.669 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

44444 443 5443	C 4 3 3 3 3 3 2 2 8 7 8 7 8 9 8 9 7 8 9 8 9 9 8 9 9 9 9 9	Result
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Sequence 27, Appl Sequence 18, Appl Sequence 18, Appl Sequence 18, Appl	Sequence 3, Appli Sequence 4, Appli Sequence 57, Appli Sequence 57, Appli Sequence 15, Appli Sequence 15, Appli Sequence 15, Appli Sequence 26, Appli Sequence 26, Appli Sequence 26, Appli Sequence 26, Appli Sequence 27, Appli Sequence 28, Appli Sequence 29, Appli Sequence 31, Appli Sequence 32, Appli Sequence 33, Appli Sequence 39, Appli Sequence 25, Appli Sequence 27, Appli Sequence 31, Appli Se	5

ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/297,181
CURRENT FILING DATE: 1999-04-26
EARLIER APPLICATION NUMBER: PCT/FR97/01921
EARLIER FILING DATE: 1997-0-27
EARLIER APPLICATION NUMBER: FR96/13176
EARLIER FILING DATE: 1996-10-29
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATENTIN VOY: 2.0
SEQ ID NO 3
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US-09-297-181-3
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GENERAL I
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Best Local Similarity
Matches 747; Conserv
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GENERAL INFORMATION:
APPLICANT: Bracco, Laurent
APPLICANT: Debussche, Laurent
TITLE OF INVENTION: ANTI-P53 SINGLE-CHAIN
FILE REFERENCE: ST96030-US
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TYPE: DNA
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Pred. No. 8.1e-205;
); Mismatches 0;
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RESULT 2
US-09-468-029-57
; Sequence 57, Applicat.
; GENERAL INFORMATION:
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; ANTI-SENSE:
US-09-468-029-57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SUQ ID NO: 57
SEQUENCE CHARACTERISTICS:
LENGTH: 1725 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
MOLECULE TYPE: 'DNA (genomic
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC COMPUTER:
OPERATING SYSTEM: PC COMPATION
SOFTWARE: PatientIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/09/468,029
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION UMBER: PCT/EP98/03791
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION UMBER: PCT/EP98/03791
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                      Matches 599;
 1318
                                                        1258
                                                                                                                 1198
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                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 8
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                cctgaagagggcctggagtggattgggatatattgatcctgagagtggtgaaactgaatat 180
                                                       AATGGAAAGAGCCTTGAGTGGATTGGAATTATTGATCCCTACTATGGTGGTACTAGCTAC 1137
TCCTCAGGTGGAGGCGGTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGGACATCGAG
              | teeteaggtggaggeggtteaggeggaaggtggetetggeggtggeggateggaeattgag
                                                                                                              ATGCAGCTCGAGACCTGACATCCGAGGACTCTGCAGTCTATTACTGTGCAAGATTCGCC
                                                                                                                                CAGGTGAAACTGCAGCAGTCTGGACCTGAGCTGGAGAAGCCTGGCGCTTCAGTGAAGATA 1017
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80.2%;
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                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 22
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 10
LENGTH: 759
TYPE: DNA
ORGANISM: human
S-09-235-073-10
                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 607; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10, Applica GENERAL INFORMATION:
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COURRENT FILING DATE: 1999-01-21
EARLIER APPLICATION NUMBER: 00/022,070
EARLIER FILING DATE: 1996-07-23
EARLIER APPLICATION NUMBER: PCT/NL97/00438
EARLIER FILING DATE: 1997-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Induction of T cell tolerance TITLE OF INVENTION: CD40/B7 antagonists FILE REFERENCE: 99-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Mark de Boer
APPLICANT: Marcel Theodorus
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atgcagctcaacagcctgacatctgaagactctgcagtctatttctgtgcaagatgggac
                                                 aatcagaagttcaagggcaaggccacattgactgtagacaaatcctccagcacagcctac
                                                              gccccgaacttccagggcaaggccactgtgactgcagacacatcctccaacacagcctac
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                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                      67.5%;
                                                                                                                                                                                                                                                         Score 504; DB 16;
Pred. No. 1.2e-134;
0; Mismatches 140;
                                                                                                                                                                                                                                                                                   Length
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US-09-468-029-59
; Sequence 59, Application
; GENERAL INFORMATION:
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                                                                                                                                                                   US-09-468-029-59
                                                                                         Query Match
Best Local S
Mätches 591
                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                         TYPE: nucleic acid
STRANDENRES: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (geno
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: UPPLICATION APPLICATION DATA:
APPLICATION NUMBER: PRILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                       LENGTH:
                                                agctgcaggagtcaggggcagaacttgtgaggtcaggggcctcagtcaatttgtcctgca
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                               AGCTACAGCAGTCTGGACCTGAGCTGGAGAAGCCTGGCGCTTCAGTGAAGATATCCTGCA
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                                                                                                         Similarity
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                                                                                           Conservative
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79.9%;
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                                                                                         0;
                                                                                         Score 501.6; DB 18
Pred. No. 8.2e-134;
0; Mismatches 149;
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Sequence 1, Application US/09297181
GENERAL INFORMATION:
APPLICANT: Bracco, Laurent
APPLICANT: Debussche, Laurent
ITILE OF INVENTION: ANTI-P53 SINGLE-CHAIN ANTIBODY FRAGMENTS AND THEIR USES
FILE REFERENCE: ST96030-US
CURRENT APPLICATION NUMBER: US/09/297,181
CURRENT FILING DATE: 1999-04-26
EARLIER APPLICATION NUMBER: PCT/FR97/01921
EARLIER APPLICATION NUMBER: PCT/FR97/01921
EARLIER FILING DATE: 1997-10-27
EARLIER FILING DATE: 1997-10-27
EARLIER FILING DATE: 1996-10-29
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 729
                                                                                                                                                                                                                                                                                        RESULT 5
US-09-297-181-1
           FEATURE:

NAME/KEY: CDS

LOCATION: (1)..(729)

-09-297-181-1
                                                             LENGTH: 729
TYPE: DNA
ORGANISM: Mus I
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                                                                                                                                   Sequence 46, Application US/08983035 GENERAL INFORMATION:
                                                                      APPLICANT: CONSEILLER, Emmanuel
APPLICANT: BRACCO, Laurent
TITLE OF INVENTION: P53 PROTEIN VARIANTS
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59
         CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc
STREET: 500 Arcola Road, Mailstop
CITY: Collegeville
STATE: PA
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Query Match
Best Local Similarity 79.2
Matches 594; Conservative
tcctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaaacagagg
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                                                                                                    gtgcaggctgáagacctggcagtttattactgcaagcaa---tcttataatctaccgacg
                                                                                                                                                                                   gtccctgatcgcttcacaggcagtggatctgggacagatttcactctcaccatcagcagt
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79.2%;
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Pred. No. 3.6e-122;
0; Mismatches 132;
                                         747
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Best Local Similarity 78.9
Matches 592; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 95/
APPLICATION NUMBER: FR 95/
FILING DATE: 19-JUL-11995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO FR9
FILING DATE: 17-JUL-11996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (610) 454-3808 NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: ST950
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                                                                                                                                                                                                                             tactatgaatacgacggctatgcttttggactactggggccaagggaccacggtcaccgtc 360
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AAGTCAAGTCAGAGCCTCTTGGA---TAGTGATGGAAAGACATATTTGAATTGGTTGTTA 528
                                                                   ATGACCCAAACTCCACTCACTTTGTCGGTTACCATTGGACAACCAGCCTCCATCTCTTGC
                                                                                                ctcacccagtctccatcttccctggctgtgtcagcaggaggaggaggtcgctatgagctgc 480
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                                                                                                                                                                                                           -TTTTACGGGGATGCTTTGGACTACTGGGGCCAAGGGACCACGGTCACCGTC
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78.9%;
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Pred. No. 3.1
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Mismatches 134;
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; NAME/KEY:
; LOCATION:
US-08-983-035-37
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US-08-983-035-37
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Query Match
Best Local Similarity
Matches 592; Conserv
                                                                                                                                                                                                                                                                TELEFAX: (610) 454-38
                                                                                                                                                                                                                                                                                              NAME: Savitzky Esq., Martin F. REGISTRATION NUMBER: 29,699
REGISTRATION NUMBER: ST950
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3816
                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Savitzky Fee
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 95/08729
FILING DATE: 19-7UL-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 'Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                           FEATURE:
                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NÜMBER: WO FR96/01111
FILING DATE: 17-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                        STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                               LENGTH:
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nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BRACCO, Laurent VARIANTS AND THERAPEUTICAL VENTION: P53 PROTEIN VARIANTS AND THERAPEUTICAL
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                  61.3%;
78.9%;
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Score 457.6; DB 13;
Pred. No. 4.1e-121;
0; Mismatches 134;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Version #1.30
                                  Length 1611;
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Indels

24;

Gaps

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Conservative

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RESULT 8
US-08-930-440-4
Sequence 4, Application US/08930480
GENERAL INFORMATION:
APPLICANT: BRACCO, Laurent
APPLICANT: SCHWEIGHOFFER, Fabien
APPLICANT: TOCQUE, Bruno
TITLE OF INVENTION: CONDITIONAL EXPRESSION SY:
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhome-Poulenc Rorer Inc.
STREET: 500 Arcola Road, Mailstop 3C43
COUNTRY: USA
COUNTRY: USA
ZIP: 19426
COUNTRY: USA
ZIP: 19426
COUNTRY: DAANNEE FORM.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
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LENGTH: 768 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: (CDNA
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APPLICATION UNMBER: US/08/930,480
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 95/03841
FILING DATE: 31-MAR-1995
PRIOR APPLICATION NUMBER: WO PCT/FR96/00477
FILING DATE: 29-MAR-1996
ATTORNEY/AGENT :NFORMATION:
REGISTRATION NUMBER: 35,135
REFERENCE/DOCKET NUMBER: 3595021-US
REFERENCE/DOCKET NUMBER: 3595021-US
TELLEPHONE: (610) 454-3808
INFORMATION FOR SEQ ID NO: 4:
SECULEMENT OF MEADURE PROTOCS:
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Best Local Similarity
Matches 589; Conserv
                601
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                                          CAGAGGCCAGGCCCAAAGCGCCTAATCTATCTGGTGTCTAAACTGGACTCTGGA
                                                                                                 AAGTCAAGTCAGAGCCTCTTGGA---TAGTGATGGAAAAACATATTTGAATTGGTTGTTA
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Pred. No. 7.6e-120;
0; Mismatches 137;
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SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 768
TYPE: DNA
TYPE: DNA
ORGANISM: Artificial Sequence
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US-08-930-480A-4
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OTHER INFORMATION:
-08-930-480a-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: TOCQUE, Bruno TITLE OF INVENTION: COND. FILE REFERENCE: ST95021-
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              ctcacccagtctccatcttccctggctgtgtcagcaggagaaaggtcgctatgagctgc
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atgacccaaactccactctctgtcggttaccattggacaaccagcctccatctcttgc
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Pred. No. 7.
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APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
FILE OF INVENTION: Antibodies Specific to KDF
FILE REFERENCE: Sequence Listings 1-34 for 381
CURRENT APPLICATION NUMBER: US/09/559,019
CURRENT FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 30
LENGTH: 714
TYPE: DNA
ORGANISM: Mouse
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Best Local Similarity
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Pred. No. 1.2e-117;
0; Mismatches 123;
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Sequence 29, Application US/0959019
GENERAL INFORMATION:
APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
ITILE OF INVENTION: Antibodies Specific to KDR and Uses Thereof
ITILE OF INVENTION: Antibodies Specific to KDR and Uses Thereof
ITILE REFERENCE: Sequence Listings 1-34 for 381-71
CURRENT APPLICATION NUMBER: US/09/559,019
CURRENT FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 29
LENGTH: 720
TYPE: DNA
ORGANISM: Mouse
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Pred. No. 3.3e-117;
0; Mismatches 124;
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APPLICANT: Yashwan
                                                                                                   TELEFAX: (617)742-7414
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1879 base pairs
                                                                                                                                                                                                                                                       FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/
FILING DATE: 07-JUNE-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                    NAME: Remillard, Jane E.
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: MX
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)27-7400
                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatientIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/364,088
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                MOLECULE TYPE: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: YAShwant M. Deo, et al.
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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NAME/KEY:
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RESULT 13
US-09-523-279-15
Sequence 15, Application US/09523279
Sequence 15, Application US/09523279
GENERAL INFORMATION:
APPLICANT: Yashwant M. Deo et al.
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS CONTITLE OF INVENTION: BINDING AGENTS
FILE REFERENCE: MXI-043CP3
CURRENT APPLICATION NUMBER: US/09/523,279
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: 09/364,088
PRIOR FILING DATE: 1999-07-30
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                                                                    COMPRISES
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; LENGTH: 1679
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; ORGANISM: Mus musculus FEATURE:
; NAME/KEY: CDS
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PRIOR FILING DATE: 1998-11
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PRIOR FILING DATE: 1996-07
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SOFTWARE: PatentIn Ver.
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APPLICANT: Yashwa
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1679 base pair:
                                                          1023
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TELEPHONE: (617)227-7400
TELEPAS: (617)227-5941
NFORMATION FOR SEQ ID NO: 15:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
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APPLICATION NUMBER:
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
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245 acctcagcagcctgacatctgaggacacaaccgtctattactgtaatgcagtcatctact 304
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LOCATION:
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                                                                            cgaacttccagggcaaggccactgtgactgcagacacatcctccaacacacgcctacctgc 244
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Pred. No. 1.5e-115;
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                                                        Best Local Similarity Matches 566; Conserv
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Best Local 9
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/468,029
                                                                                                                                                           MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                         FILING DATE:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                          TOPOLOGY:
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                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                     TYPE: nucleic acid
               1 caggtcaagctgcaggagtcaggggcagaacttgtgaggtcagggggcctcagtcaatttg 60
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CAGGTGCAGCTACAGCAGTCTGGACCTGAGCTGGAGAAGCCTGGCGCTTCAGTGAAGATA 126
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                                                         Mismatches
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Search completed: February 18, 2001, 01:29:24 Job time: 18122 sec

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Title:

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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/cgn2_6/ptodata/2/ina/GCOMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
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US-08-553-497A-27

US-08-553-497A-18

US-09-166-750-18

US-09-166-093-18

US-09-172-019-18

US-09-176-094-18

US-08-800-198-7
         US-09-296-595-7

US-08-453-497A-23

US-08-403-853-15

US-08-463-163-2

US-08-463-172A-7

US-08-279-772A-7

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Matches 584; Conserva
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                                                                                                                                                    Query .Match -
                                                                                                                                                                                                                                                                                                                            TELEFAX: 703-816-4100 INFORMATION FOR SEQ EDUNO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: ATTHUR R. CRAWFORD
REGISTRATION UNMBER: 25,327
REFERENCE/DOCKET NUMBER: 117-
TELECHOMUNICATION: TUFORMATION:
TELEPHONE: 703-816-4100
TELEFAX: 703-816-4100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon & V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                            `.l;caggtcaagctgcaggagtcaggggcagaacttgtgaggtcagggggcctcagtcaatttg
                                                                                                                                                                                                                                                          LENGTH: 810 base pairs
TYPE: nucleic acidic
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/652,507
FILING DATE: 02-Jul-1996
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Arlington
STATE: Virginia
61 tcctgcacago/tcctggcttcaacattaaagactactatatgcactgggtgaaacagagg 120
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                                        CAGGTGAAACIGCAGCAGTCTGGGGCAGAACTTGTGAGGTCAGGGACCTCAGTCAAGTTG
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1100 No. 5876691th Glebe Road, 8th Floor
                                                                                                                Conservative
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                                                                                                                                                                                                                                DNA (genomic)
                                                                                                                                59.2%;
77.9%;
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                                                                                                              Score 442.4; DB 2;
Pred. No. 2.6e-116;
0; Mismatches 136;
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                                                                                                                Indels
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RESULT 2
US-08-661-052-15
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APPLICANT: Yashwant M. Deo
APPLICANT: Yashwant M. Deo
APPLICANT: Robert Graziano
APPLICANT: Chezian Somasundaram
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: OF ANTI-FC REC
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                Sequence 15, Application Patent No. 5837243
                                                                                                                                                                                                                                                                                                                               Patent No.
        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
SOFTWARE: PatentIn Release #1.0, CURRENT *APPLICATION DATA:
                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                       STREET:
CITY: E
STATE:
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                                                                                                     COUNTRY:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 1679 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 58.7%;
Best Local Similarity 77.7%;
Matches 580; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/6
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: AIROLD, Beth E.
REGISTRATION NUMBER: 35,430
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                             aaccagggcaqtctcctaaagtgctgatctactgggcatccactagggaatctggagtcc
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                                                                            AGCCAGGCACTTCTCCCAAACTCTGGATTTATAGCACATCCAACCTGGCTTCTGGAGTCC
                                                                                                                                       CCAGCTCAAGTGT----
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Pred. No. 4.5e-115;
0; Mismatches 136;
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1476 AGGCTGAAGATGCTGCCACTTATTACTGCCAGCAACGGAGTAGTTACCCCACTCACGTTCG

1535 721

665 aggctgaagacctggcagtttattactgcaagcaa---tcttataatctaccgacgttcg

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RESULT 3
US-08-860-174A-9
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APPLICATION NUMBER: PCT/EP/96/03605

FILING DATE: August 14, 1996

INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 924 base ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9, Application Patent No. 5989830
                                                                                          LOCATION: 466..510
OTHER INFORMATION: //
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DO SOFTWARE: MS WORD, WINICURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "cDNA domains with synthetic
DESCRIPTION: linker(s)"
                                           OTHER INFORMATION: FEATURE:
                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/860,174A FILING DATE: June 16, 1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: PILLSBURY, MADISON & SUTRO LLP
STREET: 1100 New YORK Avenue, N.W.
STREET: 9th Floor, East Tower
CITY: WASHINGTON, D.C.
              NAME/KEY:
                                                                                                                                                                      OTHER INFORMATION:
                                                                                                                                                                                   NAME/KEY: mat_peptide LOCATION: 106..465
                                                                                                                                                                                                                             OTHER INFORMATION: /product-
                                                                                                                                                                                                                                                                                       CLONE:
                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: doub
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                                                                                                                                                                                                                                            NAME/KEY: sig_peptide LOCATION: 40..105
                                                                                   NAME/KEY:
KEY: misc_RNA
TION: 853..885
RINFORMATION:
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                                                                       mat_peptide
511..852
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                                                                 Sequence 17, Appli
Patent No. 5876969
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GENERAL INFORMATION: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

Fleer,

Reinhard

Fournier, Alain Guitton, Jean-Dominique

Application US/08797689

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FEATURE:

NAME/KEY: CDS.

LOCATION: 40..888

OTHER INFORMATION:
US-08-860-174A-9
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                                                                    gtgcaggctgaagacctggcagtttattactgcaagcaatcttataatctaccg---acg
                                                                                                                                                                 gtccctgatcgcttcacaggcagtggatctgggacagatttcactctcaccatcagcagt
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                               ttcggcggggcaccaagctggaaatcaaa 747
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TTCGGAGGGGGGACCAAGCTCGAGATCAAA 849
                                                                                                                                                                                                                            CAGAAGCCAGGGCAGCCTCCTAAACTGTTGATCTACTGGGCATCCACTAGGGAATCTGGA
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US-08-797-689-17
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Best Local S
Matches 559
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TITLE OF INVENTION: NOVEL BIOLOGICALLY
TITLE OF INVENTION: PREPARATION THEREOF
TITLE OF INVENTION: CONTAINING SAID POL
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-7AN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 28-7AN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Smith Ph.D. Julle K.
REGISTRATION NUMBER: P-38,619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (610) 454-3808
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 756 base pairs
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System
SOFTWARE: Word 5.1 (Pate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: FEATURE:
                +241 ctgcacctcagcagcctgacatctgaggacacaaccgtctattactgtaatgcagtcatc
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                                                                    192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/200,...
FILING DATE: 28-JUL-1994
FILING DATE: FR 92/01064
                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 559; Conserv
                                                                                                                                                                                                                                          61
                                                                                                                                                                                                         72 TCCTGCAAAGCTTCTGGCTACGCATTCAGTAGGTCTTGGATGAACTGGGTGAAGCAGAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
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ZIP: 19426
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                                                                  AATGGGAAGTTCAAGGGCAAGGCCACACTGACTGCGGACAGATCATCCAGCACAGCCTAC
                                                                                      gccccgaacttccagggcaaggccactgtgactgcagacactcctccaacacacgcctac 240
                                                                                                                                                         cctgaagagggcctggagtggattgggatatattgatcctgagagtggtgaaactgaatat 180
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                                                                                                                                     CCTGGACAGGGTCTTGAGTGGATTGGACGGATTTATCCTGGAGATGGAGATACCAAATAC 191
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                                                                                                                                                                                                                                                                                                                                                             53.5%;
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Pred. No. 2.5e-104;
0; Mismatches 170;
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                                                                251
311
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US-08-263-911-6
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Sequence 6, Application US/08263911 Patent No. 5877291
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APPLICANT: Mezes, Peter S
APPLICANT: Gourlie, Brian B
TITLE OF INVENTION: MULTIVALENT SINGLE CHAIN ANTIBODIES
NUMBER OF SEQUENCES: 23
                                                                                     PRIOR APPLICATION DATA
APPLICATION NUMBER: US 07/990,263
FILING DATE: 11-DEC-1992
ATTORNEY/AGENT INFORMATION:
NAME: "Ulmer, Duane C
REGISTRATION NUMBER: 34,941
REFERENCE/DOCKET NUMBER: C-41,014
TELECOMMUNICATION INFORMATION:
TELEPHONE: (517) 636-8104
TELEPHONE: (517) 636-8104
                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/263,911
FILING DATE: 21-JUN-1994
CLASSIFICATION: 530
                                                                         TELEPHONE: (517) 636-
INFORMATION FOR SEQ ID NO:
     SEQUENCE CHARACTERISTICS:
LENGTH: 2165 base pairs
TYPE: nucleic acid
curpannenness: double
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: Duane C. Ulmer
STREET: P.O.:Box 1967
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ZIP: 48641-1967
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 STRANDEDNESS:
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Sequence 8, Application US/08263911
Patent No. 5877291
GENERAL INFORMATION:
APPLICANT: Mezes, Peter S
APPLICANT: GOUTILE, Brian B
TITLE OF INVENTION: MULTIVALENT SI
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
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Best Local
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STREET:
CITY: N
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                    ADDRESSEE:
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Midland
            P.O.
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         Duane C. Ulmer
O. Box 1967
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70.5%;
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Pred. No. 3.
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                                                       SINGLE
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; TOPOLOGY: US-08-263-911-8
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Best Local Similarity
Matches 534; Conserv
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REFERENCE/DOCKET NUMBER: C-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (517) 636-8104
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 21-JUN-1994
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 11-DEC-1992
ATTORNEY/AGENT INFORMATION:
NAME: Ulmer, Duane C
                                                                           1674
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COMPUTER READABLE FORM:
MEDIUM TYPE: 'Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1374
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                                    533
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid STRANDEDNESS: double
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GGTACCAGCAGAAACCAGGGCAGTCTCCTAAACTGCTGATTTACTGGGCATCCGCTAGGG
                    ggtatcagcayaaaccagggcagtctcctaaagtgctgatctactgggcatccactaggg
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Pred. No. 3.3e
0; Mismatches
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US-08-553-497A-19
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                                                                                                                                                     TELEFAX: 703-243-6410
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tent No.
                                                                                                                                                                                                                                                                                          FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION NUMBER: EP 94118970.6
APPLICATION NUMBER: EP 94118970.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ROSELL, ELISABET
APPLICANT: BLASCO, FRANCESC
APPLICANT: PIULATS, JAUME
TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
NUMBER OF SEQUENCES: 30
CORRESPONDENCES: 30
                               MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                   REGISTRATION NUMBER: 33,302
REFERENCE/DOCKET NUMBER: ME
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-243-6333
                                                                                                                                                                                                                                           FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
NAME: HAMLET-KING, DIANA
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN,
STREET: 2200 CLARENDON BLVD. SUITE 1400
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                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                    ANTI-SENSE:
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                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                    CENGTH:
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   SOURCE
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BENDIG, MARY M.
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US-08-553-497A-25
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Sequence 25, Application US/08553497A Patent No. 5844093
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; LOCATION: US-08-553-497A-19
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Best Local Similarity
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TISSUE TYPE: splenocytes
IMMEDIATE SOURCE:
CLONE: 10 D 2 (single-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
698 ACACGTTCGGAGGGGGACCAAGCTGGAAATAAAA 732
                                                                                                                                                        578
                                                                                                                                                                                                                                                             542 agaaaccagggcagtctcctaaagtgctgatctactgggcatccactagggaatctggag
                                    714 -gacgttcggcgggggcaccaagctggaaatcaaa 747
                                                                             638
                                                                                                   662 tgcaggctgaagacctggcagtttattactgcaagcaatcttataaatctacc-----
                                                                                                                                                                            602 tccctgatcgcttcacaggcagtggatctgggacagatttcactctcaccatcagcagtg
                                                                                                                                                                                                                                     518 AGAAACCAGGATCCTCCCCCAGACTCCTGATTTATGACACCATCCAACCTGGCTTCTGGAG
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                                                                          CCTCAGGTGGCGGTGGCTCGGGTGGTGGGTCGGGTGGCGGCGGATCTGACATTGAGC
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                                                                                                                                                          TCCCTGTTCGCTTCAGTGGCAGTGGGTCTGGGACCTCTTACTCTCTCACAATCAGCCGAA 637
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71.7%;
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                                      Best
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                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: EP 941
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 941
FILING DATE: 02-DEC-1994
ATTORNEY_AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
                                                                                                                                                                                                                                                                                                                                 TELEFAX: 703-243-6410
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 726 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: ROSELL, E APPLICANT: BLASCO, FAPPLICANT: PIULATS, TITLE OF INVENTION:
                                                                                                                                                                                                                          MOLECULAR NO ANTI-SENSE: NO ANTI-SENSE: NO TYPE: N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUPTWARE: Patentin Release #1.0, Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION: 530
PRIOR ADDITANCE: 530
                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: ME TELECOMMUNICATION INFORMATION: TELEPHONE: 703-243-6333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: MILIEN WHITE, ZELANO & BRANIGAN, P.C.
STREET: 2200 CLARENDON BLVD. SUITE 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
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CLONE: 5 f 1 (single-chain Fv, heavy, light chain,
                                                                                                                                                                                                                                                                              MOLECULE TYPE:
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                          Local Similarity
nes 537; Conserv
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                     STRAIN: Balb/c
TISSUE TYPE: S
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                                                                                                      NAME/KEY: CDS
LOCATION: 1..726
                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
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                                                                                                                                                                                                               ORGANISM:
                                                                                                                                                                                                                                                                                            TOPOLOGY:
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1 caggtcaagctgcaggagtcaggggcagaacttgtgaggtcaggggcctcagtcaatttg 60
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ROSELL, ELISABET
BLASCO, FRANCESC
PIULATS, JAUME
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ANSELL, KEITH H
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                       Score 368.4; DB 2;
Pred. No. 2.2e-95;
0; Mismatches 186;
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US-08-553-497A-21
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                                                                                        APPLICANT: MITJANS, FRANSESC
APPLICANT: BLAŞCO, FRANCESC
APPLICANT: BLAŞCO, FRANCESC
APPLICANT: PIULATS, JAUME
TITLE OF INVENTION: ANTI-EGFR
TITLE OF INVENTION: ANTIBODIES
                                              CORRESPONDENCE ADDRESS
                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 361 tcctcaggtggaggcggttcaggcggaggtggctctggcggtggcggatcggacattgag 420
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                      ADDRESSEE:
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                                                                                                                                                                                                                                ANSELL, KEITH H.
GUSSOW, DETLEF
ADAN, JAUME
                                                                                                                                                                                                                                                                                                          BENDIG, MARY M.
                                                                                                                                                                                                                                                                                                                                   KETTLEBOROUGH,
                        MILLEN,
CLARENDON BLVD.
                                                                                             ANTI-EGFR SINGLE-CHAIN ANTIBODIES
                   WHITE,
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ZELANO & BRANIGAN, P.C.
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NFORMATION FOR SEQ ID NO:

    182 ATGAGAAATCAAGAGCAAGGCCACACTGACTGACTAGACAAATCCTCCAGCACAGCCTACA

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IMMEDIATE SOURCE:
CLONE: 3 D 3 (single-chain Fv, heavy and light chain
CLONE: plus linker)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 732 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 703-243-6333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: EP 9
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 17-MAR-
PRIOR APPLICATION DATA:
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OPERATING SYSTEM: PC-D
SOFTWARE: PATENTIN Rel-
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
                                                                                                            122
                 242
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COUNTRY:
                                                                                                                                                                                         62 cctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaaacagaggc 121
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LOCATION: 1...
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                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                                                                            tgcacctcagcagcctgacatctgaggacacaaccgtctattactgtaatgcagtcatct
                                                                                                          CCTGCAAGGCTTCCGGCTACACCTTCACCAGCCACTGGATGCACTGGGTGAAGCAGAGGG
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                                                                                                                                                                                                                                                                                                Conservative
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                             Score 364.6; DB 2; Pred. No. 2.6e-94;
                                                                                                                                                                                                                                                                                                Mismatches 184;
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RESULT 10
US-08-553-497A-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
           APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS: ADDRESSEE: MILLEN, WHITE,
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: PIULATS, JAUME
TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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ROSELL, ELISABET
BLASCO, FRANCESC
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NUMBER:
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US-08-553-497A-27
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
NAME: HAMLET-KING, DIANA
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LENGTH: 726 base pairs
TYPE: nucleic acid
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ORIGINAL SOURCE:
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TO 703-243-6410
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IOR APPLICATION DATA:
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LOCATION: 1..726
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                      aatccagtcagagtctgttcaacagtagaacccgaaagaattacttggcttggtatcagc
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                                                            TCACCCAGTCTCCAACAATCATGTCTGCATCTCCAGGGGAGAAGGTCACCATGACCTGCA
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                                                                                                                       Balb/c
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Pred. No. 2.2e-92;
0; Mismatches 192;
 ---AAGTTACATGTACTGGTACCAGC
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                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/392,338A
FILING DATE: 22-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/989,846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                        FEATURE:
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/989,846
FILING DATE: 20-NOV-1992
PRIOR APPLICATION NUMBER: US 07/796,936
FILING DATE: 25-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Filpula, David
TITLE OF INVENTION: Multivalent Antigen-Binding Proteins
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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                                                                    TYPE: nucleic acid STRANDEDNESS: both TOPOLOGY: both
                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 09
                                                                                                                                                                                                                                                               NAME: Goldstein, Jorge REGISTRATION NUMBER: 2
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ZIP: 20005
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                                     NAME/KEY:
                                                                                                                             LENGTH:
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                                                                                                                               1460 base pairs
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                 1.,1449
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Query Match Best Local Similarity

47.5%; 71.9%;

Score 354.8; DB 2 Pred. No. 1.9e-91;

2

Length 1460

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US-09-166-750-18
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                                                                                                                                                                Sequence 18, Application US/09166750 Patent No. 6025165 GENERAL INFORMATION:
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                                  APPLICANT: Wood, James F.
APPLICANT: Hardman, Karl
APPLICANT: Bird, Robert
APPLICANT: Filpula, David
APPLICANT: Rollence, Michelle
TITLE OF INVENTION: Methods for Producing Multivalent, MITTLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 23
**CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                         976
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 STREET:
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                          ADDRESSEE:
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 Washington
             E: Sterne, Kessler, G
1100 New York Avenue,
                                                                                                                                                    Whitlow, Marc
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ork Avenue, NW
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                                                                         Antigen-Binding
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CLASSLE LCATLOW
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/392,338
FILING DATE: 22-FEB-1995
PRIOR APPLICATION NUMBER: US 07/989,846
FILING DATE: 20-NOV-1992
PRIOR APPLICATION NUMBER: US 07/796,936
FILING DATE: 25-NOV-1991
PRIOR APPLICATION NUMBER: US 07/796,936
FILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A.
REGISTRATION NUMBER: 09/72,00300(
REFERENCE/DOCKET NUMBER: 09/77.00300(
TELEPHONE: (202) 371-2540
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 1460 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: US-09-166-750-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 71.>
539; Conservative
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/166,750
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                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: both
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COUNTRY: U.S:A.
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ctcacccagtctccatcttccctggctgtgtcagcaggagagaaggtcgctatgagctgc
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71.9%;
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Pred. No. 1.9e-91;
0; Mismatches 142;
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                                           GACGTCGTG
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US-09-166-093-18
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GENERAL INFORMATION:
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           TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/166,093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1036
                                                                  NAME: GOLdStein, Jorge A.
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0977.003000B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                 FILING DATE: 20-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07
                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/392,338
FILING DATE: 22-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Rollence, Michelle
TITLE OF INVENTION: Multivalent Antigen-Binding
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Whitlow, Marc
APPLICANT: Wood, James F
APPLICANT: Hardman, Karl
                                                                                                                                                               FILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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LENGTH:
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Hardman, Karl
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                             US-09-172-019-18

Sequence 18, Applicat
Patent No. 6103889
GENERAL INFORMATION:
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APPLICANT: APPLICANT: APPLICANT:

Hardman, Karl Bird, Robert

Whitlow, Marc

Application

us/09172019

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; FEATURE:
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US-09-166-093-18
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Pred. No. 1.9e-91;
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US-09-172-019-18
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Best Local Similarity
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: 0977.003000D
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
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PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: Herewith
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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FILING DATE: 25-NOV-1991
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Pred. No. 1.9e-91;
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APPLICANT: I
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               FILING DATE: 25-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A
                                                                                      APPLICATION NUMBER: US 0: FILING DATE: 20-NOV-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: Patentin Release #1.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/166,09
                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 22-FEB-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy
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TITLE OF INVENTION: Multivalent Antigen-Binding Proteins
NUMBER OF SEQUENCES: 23
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                                                    APPLICATION NUMBER: US 07/796,936 FILING DATE: 25-NOV-1991
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Hardman, Karl
Bird, Robert
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CR: 29,021
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INFORMATION FOR SEQ ID NO: 18
SEQUENCE CHARACTERISTICS:
LENGTH: 1460 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
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Best Local Similarity 71.9
Matches 539; Conservative
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
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                 ttcggcgggggcaccaagctggaaatcaaa 747
                                                        GTGAAGACTGAAGACCTGGCAGTTTATTACTGTCAGCAGTATTATAGCTATCCCCTCACG
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TTCGGTGCTGGGACCAAGCTTGTGCTGAAA 1065
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Search completed: February 17, 2001, 22:59:28 Job time: 35119 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result

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AR054409 Sequence AF226875 Synthetic AF162709 Synthetic AF169027 Synthetic	A68604 Sequence 4 AJ250763 Mus muscu AX003776 Sequence A45006 Sequence 1 AX006750 Sequence AX003768 Sequence AX0131195 Mus muscu	AJ131532 Synthetic A57272 Sequence 4 A59386 Sequence 36 A59381 Sequence 31 AB030249 Synthetic U31739 Single chai Z29480 M.musculus U44796 Synthetic AJ731533 Synthetic AJ7064776 Synthetic

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Conseiller,E., Dureuil,C., Teillaud,J.L., To
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de Genetique Moleculaire, 22 rue /
      /translation="QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQG
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/db_xref="taxon:32630"
/cell_line="PAb421"
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Sequence 4
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Bracco,L., Schweighoffer,F. and Tocque,
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Conseiller E. and Bracco, L.
P53 PROTEIN VARIANTS AND THERAPEUTIC
Patent: WO 9704092-A 36 06-FEB-1997;
RHONE POULENC ROKER SA (FR)
Other publication FR 2736915 970124.
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SLLDSDGKTYLNWLLQRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEA
EDLGVYYCWGOTHSPLTFGAGTKLELKRAAALQTRRAPAAPFPAAPAPESWPLSSS
VPSQKTYGGSYGFRLGFLHSGTAKSVTCTYSPALNKMEQQLAKTCPVQLWDSTPPPG
TRVRAMAIYKQSQHMTEVVRRCCHHERCSDSDGLAPFQHLIRVEGNLKVEYLDDRNTF
RHSVVVPYFPPEVGSDCTTIHNNYMCNSSCMGGMNRRPILTITLEDSSGNLLGRNSF
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                                                 Nakayashiki, N., Yoshikawa, K., Nakamura, K., Hanai, N., Okamoto, Nizuno, M., Wakabayashi, T., Saga, S., Yoshida, J. and Takahashi, Production of a single chain variable fragment antibody recog type III mutang epidermal growth factor receptor Unpublished (199)
3 (bases 1 to 912)
Yoshikawa, K. and Nakayashiki, N.
Direct Submission
Submitted (21-JUL-1999)
Kazuhiro Yoshikawa, Aicl
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Monoclonal antibody against the fusion junction deletion-mutant epidermal growth factor receptos Br. J. Cancer 73 (11), 1366-1372 (1996)
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Tel:81-561-62-3311(ex.2265), Fax:81-561-62-9318)
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/note="cloned from mouse hybridoma producing 3C10
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QMGRDLYDDDDKDPSSRSAAGTMEFEA"
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GLEWIGRIDPENDETKYGPIFOGGGSHMVYWTQSPLTLSVAIGGSASISKESGS
VYWGPGTTLTVSSGGGGSGGGGSGSHMVWTQSPLTLSVAIGGSASISKESGS
LLDSDGKTYLNWLLQRPGQSPKRLISLVSKLDSGVPDRFTGSGSGTDFTLRISRVEAE
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/product="3C10 single chain antibody"
/protein_id="BAA82610.1"
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3 (bases 1 to 747)
Artsaenko,O., Peisker,M., zur Nieden,U.,
Muntz,K. and Conrad,U.
Expression of a single-chain Fv antibody
creates a wilty phenotype in transgenic t
Plant. 1. 8 (5), 745-750 (1995)
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Submitted (10-JAN-1994) Artsaenko C
and Crop Plant Research, Molecular
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./note="linker sequence"
/organism="synthetic construct"
a 183 c 218 g 183 t
                                                                                                                                                                                                                                                                                                                                                                                                                         /translation="OVOLOOSGAELVRPGASYKLSCTASGFNEKEDDY.HWVKQRPEKG
LEWIGRIAPASGNVKYVPRFODKATITADTSSNTAYLLLSSLTSEDTAVYYCAR.R.S.T.
YTSLCYWGOGTTUTYVSWGGGSGGGGSGGGGSDIELTOSPPSVVVIPGESVSISCRS.
KSLLYSDGDSYLFWFLQRPGQSPQLLIYRMSNLASGVPDRFSGSGSGTSFTLRISRVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /standard_name="Vh/Vk fusion"
/note="binding abscisic acid"
/product="single chain Fv antibody"
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/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /protein_id="CAA82617
/db_xref="GI:1694843"
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/product="single chain
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Pred. No. 1.8e-144;
0; Mismatches 131;
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1 (bases 1 to 732)
Guesdon, J.-L. and Muller, B.H.
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U44796
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Synthetic construct single-chain anti-acetylaminofluorene ant
MRNA, from PCR amplified mouse VH and VL regions, partial cds
                                                                                                                                                                                                                                                                                                    Submitted (05-JAN-1996) Jean-Luc Guesdu Docteur Roux, Paris, 75015, France
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/translation="QVQLQQSGAELAKPGASVKMSCKASGYTFTTYMMHWVKQRPGQG
/translation="QVQLQQSGAELAKPGASVKMSCKASGYTFTTYMMHWVKQRPGQG
                            /transi_table=11
/product="scrv anti-acetylaminofluorene"
/protein_id="AAA86761.1"
/db_xref="GI:1177223"
                                                                                                                                                                                                                                          /organism="synthetic construct"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                         /codon_start=:
                                                                                                      /note="single-chain
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/note="light chain VJ region"
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 construct for and
g linker sequence
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Direct Submission
Submitted (16-DEC-1998) de Fromentel C.,
Cochin de Genetique Moleculaire, 22 rue 1
FRANCE
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99124402
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                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product="anti-p53 SCEV antibody (scFv11D3)"
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EXPGYALDYWGQGTTVFVSSGGGGSGGGSGGGSGTGETLTI
EXSQSLFNSRTRKNILAWYQQKPGQSBKYLLYWASTRESGVPDRFTGSGSGTDFTLTI
                                                                                                                                                                                                                                                                                                             /note=
                                                                                                                                                                                                                                                                                   /product
                                                                                                                                                                                                                                                                                                                                      /organism="Mus musculus"
/db_xref="taxon:10090"
123. .137
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                                                                                                                                                                                                                                            /organism="Mus musculus"
/db_xref="taxon:10090"
188 c 205 g 16
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/db_xref="taxon:10090"
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                                                                                                                                                                                                                                  artificial sequence.

1 (bases 1 to 738)

Tewari,D., Goldstein,S., Notkins,A.L. and Single chain antibody to HIV-1 pl7

J. Immunol. (1998) In press

2 (bases:1 to 738)
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                                                                    /organism="synthetic construct"
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./db_xref="ARTCC:HB8975"
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HIV-1 p17 (ATCC HB8975)"
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/transl_table=11
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SLLGSDGKTFLNWLLQRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKISRVEA
EDLGVYYCWQGTHLPQTFGGGTKLEIK"
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               CCACCCAAACTCCTCATCAAGTATGCATCCAACCTAGAATCTGGGGTCCCTGCCAGGTTC
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Aspergillus clavatus

Eukaryota; Fungi; Ascomycota; Euascomycete

Eurotiales; Trichocomaceae; mitosporic Tri

(bases 1 to 1314)

Mele, A., De, S.R., Parente, D. and Colnaghi,

RECOMBINANT RIBOSOMAL INHIBITOR PROTEIN (F
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Patent: WO 9749726-A 31-DEC-1997;
MINISTERO UNI RICERCA SCIENT E (IT)
Other publication IT F1960155 19971229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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/strain="IFO8605"
/db_xref="taxon:5057"
1 360 c 359 g 2
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Pred. No. 1.8e-129;
0; Mismatches 153;
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Medizinische Hochschule
Hannover, GERMANY
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and their application in xenotransplantation
MO1. Immunol. 38, 1235-1247 (1999)
2 (bases 1 to 759)
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synthetic construct
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ScFv; variable region.
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                                                        Similarity
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1 (bases 1 to 759)
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                                             Conservative
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                                                                                                                                                                                        /organism="Mus musculus"
/db_xref="taxon:10090"
376. .420
                                                                                                                                                                                                                                                                                         /product-"anti-guinea pig C5 ScFv antibody"
/protein_id="CAB60136.1"
/protein_id="CAB60136.1"
/db_xref="G1:6272279"
/translation="QPAMAEVQLQQSGAALARPGASVKLSCRASGYTFSSYWLQWVKQ
/translation="QPAMAEVQLQQSGAALARPGASVKLSCRASGYTFSSYWLQWVKQ
RPGGGLEWIGARYFGGDTRYNQKFKGKATMTADKSSTTAYMQLTILSSEDSAVYYCA
RYGDYLYGMDYWGQGTSVTVSSGGGGSGGGGSGGGSDVVMTQTPLSLPVSLGDQAS
ISCRSSGSLVHSSNGNTYLHWYLQKPGQSPKLLIXKVSNRFSGVPDRFSGSGSGTDFTL
KISKVERAEDLGVYFCSQSTHVPWTFGGGTKLEIKR"
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420. .7
                                                                                                              /organism="Mus musculus"
/db_xref="taxon:10090"
183 c 213 g 18
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/db_xref="taxon:10090"
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/transl_table=11
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/db_xref="taxon:32630"
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                                           Score 446.4; DB 12;
Pred. No. 1.1e-127;
0; Mismatches 156;
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                                                                                                                         Method of identifying binding sit of binding to an epitope patent: WO 9925818-A 27 MAY-1999;
                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalla; Euthería; Rodentía;
1 (bases: 1 to 753)
Kufer,P and Raum,T.
                                                                                                     KUFER PETER (DE); RAUM TOBIAS
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/organism="Mus sp."
/db_xref="taxon:10095"
188 c 215 g
                                                                               Location/Qualifiers
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1 (bases 1 to 810)
Chester,K.A., Hawkins,R.E. and Begent,R. ANTIBODY AGAINST CARCINOEMBRYONIC ANTIGE Patent: WO 9515341-A 1 08-JUN-1995;
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Other publication AU 1194795
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                                              ATCTAGTCAGAGCCTTGTACACAGTAATGGAAACACCTATTTACATTGGTACCTGCAGAA 1364
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17 from Patent
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/db_xref="taxon:32630"
/note="artificial sequence"
/note="artificial sequence"
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                tgctggcaccuagctggaaatcaaa
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completed: February 35030 sec 17, 2001, 22:57:49

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Listing first 45 summaries
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gb_gss26: *
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gb_gss28: *
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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189 189 188 187 187	210 2204 2004 2004 198 198 198 199 199 199 199 199 199 199	A, A, A, A, A, A, A, A, A, A, A, A, A, A	S
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ALIGNMENTS

ACCESSION VERSION KEYWORDS SOURCE BE306420 *, 1057 bp mRNA EST 13-JUL-2000 601103347F1 NCI_CGAP_Lu29 Mus musculus cDNA clone IMAGE:3495755 5', BE306420 BE306420 1 GI:9159604 EST. house mouse.

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RESULT
BE307894
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mRNA sequence.
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                                      BE307894 793 bp mRNA
601096548F1 NCI_CGAP_Mam5 Mus
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Plate: LLAM8546 row: n column:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      National Institutes of Health, Mammalian Gene Collection (MGC) 
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNE at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammmalla; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: Robert_Strausberg@nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NIH-MGC http:,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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289 c 273 g 221 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /strain="CZECH II (feral)"
/db_xref="taxon:10090"
/clone="IMAGE:3495755"
/clone=lib="NCI_CGAP_Lu29"
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Pred. No. 2.3e
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KEYWORDS
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                                                                                                                                       CTCACGTTCGGTGCTGGGACCAAGCTGGAGCTGAAA
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AW405772 471 bp mRNA EST 16-FEB-2000 UI-HF-BLJ-abp-c-06-0-UI.rl NIH_MGC_37 Homo sapiens cDNA clone
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1 (bases 1 to 793)
NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.
National Institutes of Health, Mammalian
Unpublished (1999)
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM8534 row: p column: 12
High quality sequence stop: 514.
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Eukaryota; Metazoa;
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'Site_2: NotI; Cloned unidirectionally. Primer: Oligo d'
Library constructed by Life Technologies. Investigators
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         providing samples: Lothar Hennighausen/Robin Humphreys,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross tissue"
/dev_stage="7 months"
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/strain="C57/B6"
/db_xref="taxon:10090"
/clone="IMAGE:3491195"
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Pred. No. 4.7e-60;
0; Mismatches 63;
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                                                                                                                                                                     tctaaactggactctggagtccctgacaggttcactggcagtggatcagggacagatttc 621
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                                                                                                                                                                                                                        TATTTGTATTGGTACCTGCAGAAGCCAGGCCAGTCTCCACAGCTCCTGATCTATGAAGTT 194
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Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

CDNA Library Preparation: M.B. Soares Lab

CDNA Library Arrayed by: M.B. Soares Lab

DNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.linl.gov/bbrp/image/image.html

Seq primer: M13 Forward.
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Tel: (301) 496-1550
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Constructed from size fractionated cytoplasmic mRNA (1.5-2.5kb). Directionally cloned. Cells provided by Low M. Staudt, Ph.D. Library preparation by Maria de Fatima Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

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/cell_type="action"
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/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
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/cell_line="MGC85"
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/db_xref="taxon:9606"
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Pred. No. 2.1e-59;
Pred. No. 2.2e-59;
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355
                 682 acacattctccgctcacgttcggtgctggcaccaagctggaaatcaaa 729
                                                                                          622 acactgaaaatcaacagagtggaggctgaggattttggggagtttattattgctggcaaggt 681
                                                                                                                                                 235 TCCAACCGGTTCTCTGGAGTGCCAGATAGGTTCAGTGGCAGCGGGTCAGGGACAGATTTC
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ATACAGCTTCCTCGCACGTTCGGCCAAGGGACCAAGGTGGAAATCAAA
                                                                        Eco RI site shown at the beginning of the sequence. Tissue procurement: Louis M. Staudt, M.D., Ph.D. cDNA Library Preparation: M.B. Soares Lab cDNA Library Arrayed by: M.B. Soares Lab DNA Sequencing by: M.B. Soares Lab Clone distribution: MGC clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: www-bio.i.nl.gov/bbrp/image/image.html
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1. (bases 1 to 614)
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UI-HF-BLO-acb-c-06-0-UI.rl NIH_MGC_37 Homo
IMAGE:3058546 5', mRNA sequence.
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Tel: (301) 496-1550
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National Institutes of Health, Mammalian
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//note="Vector: pT773-PG: Site_1: Not1; Site_2: Eco RI;
/note="Vector: pT773-PG: Site_1: Not1; Site_2: Eco RI;
/constructed from size Parattionated cytoplasmic mRNA
(1.5-2.5kb). Directionally cloned. Cells provided by Lo:
// Y. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

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/db_xref="taxon:9606"
/clone="IMAGE:3058546"
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/cell_line="MGC85"
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/tissue_type="lymph"
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TITLE
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Best Local
                  621 cacactgaaaatcaacagagtggaggctgaggatttgggagtttattattgctggcaagg 680
                                                                                                                                                381 cggtggcggatcggatgttttgatgacccaaactccactttgtcactttgtccggttaccatttgg 440
                                                                                                                                                                                                                           441 acaaccagcctccatctcttgcaagtcaagtcagagcctcttggatagtgatggaaagac
                                                                                                                                                                                              63
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nes 274; Conserv
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                                                          CTATTTGGATTGGTACCTGCAGAAGCCAGGGCAGTCTCCACAGCTCCTGATCTATTTGGG
                                                                                                                                                                                                                                                             CGAGGCCAGTGGGGATATTGTGATGACTCAGTCTCCACTCTCCCTGCCCCGTCACCCCTGG 62
                                                                                                                                                                                              AGAGCCGGCCTCCATCTCCTGCAGGTCTAGTCAGAGCCTCCTGCATAGTAATGGATACAA 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AW406883 454 bp mRNA EST 16-FEB-2000 UI-HF-BLO-adg-h-03-0-UI.rl NIH_MCC_37 Homo sapiens cDNA clone IMAGE:3061493 5', mRNA sequence.

AW406883 AW406883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seq primer: M13 Forward.
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nmalia; Eutheria; (bases 1 to 454)
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/clone_lib="NIH_MGC_37"
/tissue_type="lymph"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="DH10B (LTI)"
note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
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/db_xref="taxon:9606"
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78.5%;
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_line="MGC85"
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                                                                                                                                                                                                                                                                                                                                           Score 229; DB 90;
Pred. No. 2.9e-58;
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AUTHORS
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                                                                                                                                                                                                                                                                                      394 gatgttttgatgacccaaactccactcactttgtcggttaccattggacaaccagcctcc 453
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                                                                                                               126 TTTCAGCAGAGGCCAGGCCAATCTCCAATGAGCCTAATTTATAAGGTTTCTAACCGGGAC
                                                                                                                                   514 .ttgttacaga;gccaggccagtctccaaagcgcctaatctaggtgtctaaactggac
634 aacagagtggaggctgaggatttgggagtttattattgctggcaaggtacacattctccg
                                                      574 totggagtccatgacaggttcactggcagtggatcagggacagatttcacactgaaaatc
                                     186 TCTGGGGTCCCAGACAGATTCAGCGGCAGTGGGTCAGGCACTGACTTCACACTGAAAATC
                                                                                                                                                                                        66 ATCTCCTGCAGGTCTAGTCAAAGCCTCGCACACAGTGATGGAAACACCTACTTGAATTGG
                                                                                                                                                                                                                                                                 6 GGTGTAGGGATGACTCAGTCTCCACTCTCCCTGCCCGTCACCCTTGGACAGCCGGCCTCC
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                                                                                                                                                                                                                                                                                                                                         268;
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NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eco RI site shown at the beginning of Tissue Procurement: Louis M. Staudt,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999
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cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
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/clone="IMAGE:3060080"
/clone_lib="NIH_MGC_37"
/tissue_type="lymph"
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/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2:
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/cell_line="MGC85"
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db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                           Score 227.2; DB 90; Pred. No. 1e-57;
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AUTHORS
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                                                                                                                  ggatgttttgatgacccaaactccactcactttgtcggttaccattggacaaccagcctc
                                 CATCTCCTGCAGGTCTAGTCAGAGCCTCCTGCATAGTAATGGATACAACTATTTGGATTG 137
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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IMAGE:809878 5' similar to gb:X72467
REGION (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hillier, L., Allen, M. Kucaba, T., Lacy, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA464313.1 GI:2189197
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Seq primer: -28ml3 rev2_ET from Amersham
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314 286 1810
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                                                                                                                                                                                                                                                                                                                    the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M Fatima Bonaldo "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:809878"
/clone_lib="Soares ovary tumor NbHOT"
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/db_xref="GDB:6039712"
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                                                                                                                                                                                                                                                                                                                                                                                                                                   'tissue_type="ovarian tumor"
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                                                                                                                                                          Score 223.4; DB 7; Pred. No. 1.3e-56; 0; Mismatches 71;
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Tan,F., Theising,B., White,Y., Wylie
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7 IG KAPPA CHAIN PRECURSOR V-II
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Query Match Best Local S Matches 266

Similarity

30.6%;

Score 223.4; DB Pred. No. 1.4e-56 Mismatches

.4e-56;

24;

71;

Indels Length

0

Gaps

0

Conservative

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BASE COUNT
ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            318 ATTCACTTTCGGCCCTGGGACCAAAGTGGATATCAAA 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            198 CTCCGGGGTCCCTGACAGGTTCAGTGGCAGTGGATCAGGCACAGATTTTACACTGAAAAAT 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cancer Genome Anatomy Project (CGAP), Tumor Gene Index Unpublished (1997)
Other_ESTs: zu56d02.sl
Contact: Robert Strausberg, Ph.D.
Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMAGE Consortium (info@image.llnl.gov) for further informa This read is a RESEQUENCE of a previously sequenced human Original clone citation: WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: (301) *** ****
Email: Robert_Strausberg@nih.gov

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This clone is available royalty-free through LLNL; contact the

This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AI734035 ' 425 bp' mRNA EST 14-JUN-1999 zu56d02.y5 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:741987.5' similar to gb:X72467 IG KAPPA CHAIN PRECURSOR V-II
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This read has been verified (found to hit its original self in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI/NIDR CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        orientation)
                       (Pharmacra, M.Fatima Bonaldo. "
M.Fatima Bonaldo. "
114 c 111 g
                                                                 double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and
                                                                                                                                                             (Pharmacia). Library
                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="GDB:5942254"
/db_xref="taxon:9606"
/clone="IMAGE:741987"
                                                                                                                                                                                                                                                        /tissue_type="ovarian tumor"
/lab_host="DH10B (ampicillin
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                       /sex="Female"
                                                                                                                                                                                                                                                                                                                        /clone_lib="Soares ovary tumor NbHOT"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   White,Y., Wylie,T., Waterst WashU-NCI human EST Project
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                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Wilson RK
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314 286 1810
   /organism="Homo sapiens"
/db_xref="GDB:5942254"
/db_xref="taxon:9606"
/clone="IMAGE:741987"
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x 8501, St. 1
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KAPPA CHAIN PRECURSOR
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AW405725.
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UI-HF-BLO-abv-f-05-0-UI.rl NHLMGC_37 Homo
IMAGE:3057944 5', mRNA sequence
AW405725:
                                                                                                                                                                                                                                                 Email: Robert_Strausberg@nih.gov
ECO RI site shown at the beginning of the se
Tissue Procurement: Louis M. Staudt, M.D.,
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution.
                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
Contact: Robert Stra
Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                   NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 488)
                                                                                                                                                                                                   Seq
                                                                                                                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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primer: M13 Forward.
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/cell_type="germinal center B cells"
/cell_line="MGC85"
/lab_host="DH10B (ITI)"
/lab_host="DH10B (ITI)"
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco
Constructed from size fractionated cytoplasmic mRNA
                                                                              /clone="IMAGE: 3057944"
/clone_lib="NIH_MGC_37"
/tissue_type="lymph"
                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                  Location/Qualifiers
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78.9%;
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                            This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scriptts/gethtm12.pl?tl=&t2=RC6-ET0073-210 600-011-F01&t3=2000-06-21&t4=1) see primer: puc 18 forward the primer: puc 18 forward the primer:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleo Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (Dase: 1 to 456)

National: Institutes of Health, Mammalian Gene Collection (Mupublished (1999)

Contact: Robert Strausberg, Ph.D.

Tel: (301) 496-1550

Email: Robert Strausberg@nih.gov

Eco Rf site shown at the beginning of the sequence.

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

CDNA Library Arrayed by: M.B. Soares Lab

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LINL at:

www-bio.llnl.gov/bbrp/image/image.html
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a 121 c 118 g 107 t 1 others
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                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 488)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.
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yl148q05.rl Soares breast 3NbHBst Homo
IMAGE:161528 5' similar to gb:X72467 I
REGION (HUMAN);, mRNA sequence.
                         Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
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                                                                                         Contact: Wilson RK
                                                                                                                  Unpublished (1995)
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/note="Vector: pT/73-9c; Site_1: Not1; Site_2: Eco RI;
/note="Vector: pT/73-9c; Site_1: Not1; Site_2: Eco RI;
/note="Vector pt/73-9c; Site_1: Not1; Site_2: Eco RI;
/note="Vector pt/73-9c; Site_1: Not1; Site_2: Eco RI;
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/note="Vector pt/73-9c; Site_1: Not1; Site_2: Eco RI;
/note="Vector pt/73-9c; Site_1: Not1; Site_1
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/clone_lib="NIH_MGC_37"
/tissue_type="lymph"
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/cell_line="MGC85"
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/db_xref="taxon:9606"
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0; Mismatches 8:
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601092470F1 NCI_CGAP_Mam5
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Mus muscalus
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Insert Size: 1071
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/clone="IMAGE:161528"
/clone_lib="Soares breast 3NbHBst"
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/db_xref="GDB:575685"
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AW406473 353 bp mRNA UI-HF-BLO-acr-e-08-0-UI-II NIH_MGC_37 IMAGE:3060182 5', mRNA sequence.
AW406473.1 GI:6925530
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Tel: (301) 496-1550
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: Robert_Strausberg@nih.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sal. Site_2: NotI; Cloned unidirectionally. Primer: Oligo di Library constructed by Life Technologies. Investigators providing samples: Lothar Hennighausen/Robin Humphreys,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="IMAGE:3487241"
/clone_lib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross tissue"
/dev_stage="7 months"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism-"Mus musculus"
/strain-"C57/B6"
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Pred. No. 7.2
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Eco RI site shown at the beginning of the sequence.

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

cDNA Library Preparation: M.B. Soares Lab

cDNA Library Arrayed by: M.B. Soares Lab

CNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.lnl.gov/bbp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          www-bio.lini.gov/bbrp/image/image.html
Seq primer: M13 Forward.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
Contact: Robert Stri
Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 353)
NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/
National Institutes of Health, Mammalian
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Mammalia, Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Constructed from size fractionated cytoplasmic mRNA (1.5-2.5kb). Directionally cloned. Cells provided by M. Staudt, Ph.D. Library preparation by Maria de Fai Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

93 c 93 g 94 t
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/db_xref="taxon:9606"
/clone="IMAGE:3060182"
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/tissue_type="lymph"
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/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2:
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Search completed: February Job time: 13895 sec

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729
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Copyright (c) 1993 - 2000 Comp
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Coding sequence fo
Anti-human protein
Human p53 protein
Plasmid pJG718 enc
DNA od SCFV D3M wh
Mouse scrV fragmen
MFE-23 antibody co
Bispecific anti-ze
Mouse scrV fragmen
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Fusion gene sequen	V00687	19	1797	7.	49.	4
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pscFvNQ11 encoding	021098	13	747	8	51.	2
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cDNA encoding an a	X86614	20	2478	8	52.	ö
Antibody-Firefly 1	V23580	19	w	8	54.	9
Firefly luciferase	V23579	19	732	8	54.	8
Bispecific tetrava	X01651	20	1725		56.	37
Mouse scFV fragmen	X77243	20	744	9	57.	8
scFV fra	X77248	20	744	9.	60.	5
H and L chain FV r	Q35955	14	778	9.	61.	34
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scFV frag	X77247	20	744		68.	
Fusion protein pNG	V72064	20	0	:	77.	
smid	V72059	20	2019	۲.	77.	
sion pro	V72072	20	864	:	77.	
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Single chain anti-	T58130	18	S)		25.	ω

ALIGNMENTS

RESULT V36236

V36236;

V36236 standard; DNA; 729 BP

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(RHON ) RHORE-POULENC RORER
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Restoring p53-dspendent trans-activating activity to cell containing mutant p53 - by delivering single-chain antibody specific for the
                                                                                                                                                           Debussche L,
                                                                                                                                                                                                                                                         29-OCT-1996; 96FR-0013176.
                                                                                                                                                                                                                                                                                                                27-OCT-1997;
                                                                             WPI; 1998-272140/24.
P-PSDB; W60769.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumour-suppressing activity;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                           The invention relates to novel chimaeric, bispecific proteins which CC comprise: (a) a DNA binding domain and (b) a domain which binds a CC trans-activator (TA), trans-repressor (TR) or their complexes, which are CC trans-teristic of a physiological or physiopathological state. The novel CC chimaeric, bispecific proteins allow expression of a therapeutic protein CC (e.g. diphtheria or Pseudomonas toxins, thymidine kinase, single chain CC antibodies) to be regulated in response to particular conditions. CT The Ta- or TR-binding domain may be a antibody such as a single chain CC in the p53 protein. The sequence is constructed from the gene encoding CC to the p53 protein. The sequence is constructed from the gene encoding CC the light and heavy chain variable regions of the antibody produced by CC hybridoma pAb421. The 340 and 325 by sequences encoding the heavy and CC light chain variable regions respectively are amplified by PCR and CC included to the specification), amino acids 9-241 represent the linked CC whilnker-V1 sequence. The protein also contains an myc epitope tag crepresented by amino acids 256-266.
                                                               Query
Best I
                                            Best Local Sir
Matches 722;
                                                                                                                    -Sequence
     | Caggtgcagctgcagcagtctgggggcagagcttgtgaggtcagggggcctcagtcaagttg
                                                                                                                                                                                                                                                                                                                                                                                                                            Conditional gene expression system triggered by e.g. infection hyper-proliferation - comprises novel bi:specific proteins have DNA-binding domain and second domain specific for trans-activates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diphtheria; Pseudomonas; toxin; thymidine kinase; pathogen; HIV Tat; papilloma virus; E6/E7; Epstein hyperproliferation; p53; tumour; oligomerisation;
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single chain antibody;
n-Barr virus; EBNA;
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substitution; replacement; transactivation; viral protein VP16;
anti-oncogene; hyperproliferation; cancer; restenosis; SCFV;
tumour suppression; apoptosis; single chain antibody variable do
                                     17-JUL-1996;
                                                                                                                                         Synthetic
                                                                                                                                                           Mammalia
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CC Claimed variants of protein p53 have at least part of the CC coligomerisation domain deleted and replaced by a leucine zipper domain. The mutants preferably also have at least part of the p53 cransactivation domain (amino acids 1-74) deleted and replaced by the transactivating domain (TD) from herpes simplex virus viral crowder of the transactivating domain (TD) from herpes simplex virus viral crowder of the transactivating domain (TD) from herpes simplex virus viral crowder of the transactivation of the protein domain able to bind selectively to a transactivator, especially a single-chain cantibody variable domain (SCFV). The present sequence represents the NcoI-NotI fragment of cDNA coding for the anti-p53 single-chain coding for the sequence represents the NcoI-NotI fragment of cDNA coding for the anti-p53 single-chain coding sequence to the indicated) and no termination codon. The p53 variants are more active and more stable tumour suppressors and capoptosis-inducing agents than wild-type p53 and are active where the wild-type protein is not, i.e. they are not inactivated by companion of coding the sequence or oncogenic mutants, nor by other cellular corrections (because the leucine zipper domain prevents formation of codinactive mixed oligomers).
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Claimed variants of protein p53 have at least part of the oligomerisation domain deleted and replaced by a leucine zipper domain. The mutants preferably also have at least part of the p53 transactivation domain (amino acids 1-74) deleted and replaced by the transactivating domain (TD) from herpes simplex virus viral protein vP16 (amino acids 411-490) or by a protein domain able to bind selectively to a transactivator, especially a single-chain
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substitution; replacement; transactivation; viral protot
anti-oncogene; hyperproliferation; cancer; restenosis;
tumour suppression; apoptosis; single chain antibody v
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98.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 716.2; Pred. No. 1.5e-196; O; Mismatches 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              341 T; 0 other;
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RESULT X58936 ID X5

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                                                                                                                                                                                                                                                                                                                                                                                                     derived growth factor receptor (PDFR-R TM). pJG718 expression vector was constructed by inserting a DNA segment encoding the A77 serv in frame with the PDGF-R TM sequence in plasmid vector pDisplay. Murine tumour cells transformed to express A77-TM were able to bind a soluble form of the Fc alpha receptor, and the bound receptor was able to engage IgA molecules. This is an example of cells of the
                                                                                                                                                                                                                                               are used to treat cancer and infectious diseases or used as vaccines. The method allows for killing of target cells without targeting any particular antigen on the cell. This is advantages since many tumour cells and other target cells do not have define
                                                                                                                                                                                                                                                                                                                                        component which binds to an FC receptor of an effector cell transformed cell is targeted to an effector cell via the FC component, and can be used as a vehicle to increase an eff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDS
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                                                                                                                                                                                                                                                                                                             cell mediated immune response, such as cell lysis and phagocytosis, against an antigen associated with the cell. The transformed cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This is the DNA sequence of a portion of vector pJG718 that codes for a fusion protein (see v06273) comprising murine anti-rc alpha receptor antibody A77 srv and the transmembrane domain of platelet derived growth factor receptor (PDFR-R TM). pJG718 expression
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                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cells expressing anti-Fc receptor binding components
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                                                                                                                                                                                                                                                                                                                                                                                         invention that have been transformed to express on their surface
246 cctgcaaggcttctggttattcattcactgactacatcatattttgggtgaagcagagcc:305
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              cctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaagcagaggc 121
                                                        agatccagctgcagcagactggacctgagctggtgaagcctggggcttcagtgaagatat 245
                                                                         aggtgcagctgcagcagtctggggcagagcttgtgaggtcaggggcctcagtcaagttgt
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81.2%;
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                                                                                                                                                                                                 273 C; 310 G;
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Pred. No. 7.2e-136;
0; Mismatches 120;
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                                                                               07-MAY-1998
(RHON ) RHONE-POULENC RORER SA
                                                                                                                                    Mus sp
                                                                                                                                                                         Single chain antibody; ScFv D3M; mouse; p53 protein; regulatory domain; p53 mutant; H273; W248; G281; p53-dependent trans-activating activity; restoration; tumour-suppressing activity; tumour ceil; treatment;
                                                                                                        WO9818825-A1
                                                                                                                                                            tumour-suppressing activity;
hyper-proliferation; cancer;
                                                                                                                                                                                                                                              DNA od ScFv D3M;which binds to mutant p53 proteins
                                                                                                                                                                                                                                                                            8661, das-80
                                                                                                                                                                                                                                                                                                      V36237;
                         29-0CT-1996;
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Best Local Similarity
Matches 594; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Restoring p53-dependent trans-activating activity mutant p53 - by delivering single-chain antibody smutant, particularly for treatment of tumours
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           This nucleotide sequence encodes a fusion protein comrising of the Mgr6 single chain antibody (with the heavy and light chain joined by a linker), fised to the clavin protein. Clavin is an inhibitor protein, and furctions by inactivating the ribosomes. Clavin or it conjugates are useful as anticancer and/or antiviral agents. The recombinant ribosomal inhibitor protein (RIP), Mgr6-clavin (a conjugication) is used as an immunoconjugate, and the complex and clavin of clavin is used as an immunoconjugate, and the complex and clavin alone are capable of inhibiting ribosomal activity.
                                                                                                                               Claim 3;
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Best Local Similarity
Matches 573; Conserv
                       Type II collagen; assay; antibody; rheumatoid arthritis; osteoarthritis; diagnosis; monoclonal antibody 5109; MAb; antibody engineering; scfv; single chain antibody; ss.
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comprising the heavy and light chain antibody (see Y06132) comprising the heavy and light chain variable regions of a novel murine monoclonal antibody (MAb), 5109, that is specific for type II collagen necepitopes. The sequence was assembled using a splicing by overlap extension (SOEing) method in pCANTAB6. E. coli containing the 9A4 scfv antibody plasmid is deposited as ATCC 98594. A claimed method for monitoring biological media for protein fragments, especially those generated by collagenase cleavage of articular cartilage or type II collagen, involves: contacting the biological media with a capture antibody that is active against the collagen peptides given in Y06122 and Y06123; contacting the biological media with a detection antibody that is active against the collagen fragments given in Y06124 and Y06125; and detection the collagen fragments bound to the capture and detection antibody, and then the capture antibody. The method is useful for monitoring collagen fragments in biological media, and for assessing the breakdown of type II collagen by collagenase in normal and pathological (rheumatold arthritis, osteoarthritis) conditions.

MAb 5109, and can be used to detect type II collagen fragments in body fluids independent of the method used to generate the collagen
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Sequence 753

BP; 171 A;

188 C;

215 G; 179 T; 0

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or multivalent polypeptides that comprise antibody binding sites capable of efficiently binding to the corresponding antigen. The polypeptides or antibodies identified by the method are useful therapeutically and diagnostically, for e.g. cancer and autoimmune diseases. ScFv-antibody fragments that bind independently of their position within bifunctional single-chain fusion proteins can be isolated from combinatorial antibody libraries using the new in vitro method. Sequences X77240-248 represent DNA sequences encoding mouse scFV fragments.
                                                                                                                                                                                                The invention relates to a method of identifying binding site domains (BSD) that retain the capacity of binding to a predetermined epitope when positioned C-terminal of at least one further domain in a recombinant bior multivalent polypeptide. The method comprises (a) testing a panel of BSD displayed on the surface of a biological display system as part of a fusion protein for binding to a predetermined epitope, where the fusion protein comprises an additional domain positioned N-terminal of the BSD and an amino acid sequence that mediates anchoring of the fusion protein to the surface of the display system; and (b) identifying a BSD that binds to the predetermined epitope. The method is useful to identify bi-
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therapy; d
       This sequence represents the coding sequence for the MFE-23 antibody. The encoded protein is an antibody against carcinoembryonic antigen (CEA). CEA is a marker antigen for cancer imaging and therapy. The MFE-23 antibody sequence was obtained using phage technology. In this process, mice were immunised with CEA. The antibody variable region genes obtained from these mice were then amplified from cDNA and cloned
                                                                                  Claim 3;
                                                                                                                                                                   Begent RHJ,
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diabody;
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                                                                                  Page 48-49; 72pp;
                                                                                                                carcinoembryonic antigen
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                                               ~721
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709 ggcaccaagctggaaatcaaa
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76.7%;
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Pred. No. 5.
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Mismatches 143:

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                                                                                                                                                                                                                                                                                                  an antibody which specifically interacts with the extracellular domain of CC the human zeta-chain. The antibody whose CDR of a variable region is encoded by (I), is obtained by immunising a rat with jurkat cells and CC encoded by (I), is obtained by immunising a rat with jurkat cells and CC extractions of the comprising a carrier molecule and a peptide having 11 N-terminal amino acids of a rat zeta-chain. The CC enti-zeta-chain antibody is useful for the treatment and prevention of CC autoimmune diseases, immune deficiencies, T-cell malignancies, in fectious diseases, immune deficiencies, T-cell malignancies, CC or viral infections. The antibody, and fragments of it, can be useful for the enhancement or suppression of NK-cell dependent immunity or for the CC treatment of NK-cell derived malignancies. It can also be useful for the CC treatment of the cell derived malignancies. It can also be useful for the CC treatment of their precursors. The present sequence encodes a Dispecific anti-zeta-chain/anti-EpcAM antibody, from an example from
                                                                                                                                                                           Query Match
Best Local Similarity
Matches 561; Conserv
                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 9; Page 74; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New oligonucleotide, polypeptide, antibody useful for treating autoimmune disease, immune deficiencies, T-cell malignancies and infectious diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reiter
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caaggcttctggctacaccttcacaaactatggtttaagctgggtgaagcagaggcctgg 944
                                                               cacagettetggetteaacattaaagaetaetatatgeaetgggtgaageagaggeetga 125
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P-PSDB; Y17958
               WPI; 1999-338004/28
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une disease; scFv-antibody; single-chain Fv; mouse; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                          gccaggccagtctccaaagcgcctaatctatctggtgtctaaactggactctggagtccc
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75.0%;
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Pred. No. 4.5e-114;
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                            A mammalian expression construct encoding a single chain antibody having binding specificity for Fc gamma RI, derived from the humanised anti-Fc gamma RI monoclonal antibody H22 was prepared (see T58129). A bispecific single chain polypeptide was produced by fusing the H22 SCFv to an anti-carcinoembryonic antigen (CEA) antibody. The H22-anti-CEA fusion protein was shown to bind both Fc gamma RI and CEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Humanised antibody; anti-Fc receptor; H22; bifunctional; bispecific; fusion protein; chimera; carcinoembryonic antigen; CEA; MFE-23; ds.
Sequence 1679 BP; 404 A; 440 C;
                                                                                                                                                    Example 8; Fig 40; 115pp;
                                                                                                                                                                                               Recombinant, multi-specific anti-Fc receptor antibody molecules also comprise an anti-target portion, used for the treatment of
                                                                                                                                                                                                  also
                                                                                                                                                                                                                                                           WPI; 1997-052242/05
                                                                                                                                                                                                                                                                                           Deo YM,
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Query Match
Best Local Similarity

58.3%; 76.4%;

Score 425.2; DB 18; Pred. No. 7.3e-113;

Length 1679;

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                                                           Multispecific single chain antibody; antibody H22; tumour cell; antibody-dependent cellular cytotoxicity; ADCC; HER 2/neu; infecepidermal growth factor receptor; breast cancer; ovarian cancer;
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                                                                                                                                                                                                                                                                             This sequence encodes the construct 321, which is a fusion between an anti-CEA antibody and a multispecific single chain antibody designated H22. The antibody can be used in the method of the invention for inducing antibody-dependent cellular cytotoxicity (ADCC) against a tumour cell which is characterised by overexpression of HER 2/neu or epidermal growth factor receptor (EGFR), comprises contacting the tumour cell with a multispecific protein molecule (preferably a single chain antibody) comprising: (a) an anti-Fc receptor antibody or an antigen binding comprising: (b) a portion which binds to HER 2/neu, and (c) a portion which binds to TERP material can be accorded.
                                                                                                                                                                                                                       fragment; (b) a portion which binds to HER 2/neu; and (c) a portion which binds to EGFR. The method can be used for treating cancers especially breast cancer or ovarian cancer. The multispecific antibody can also be administered prophylactically to vaccinate a subject against infection by a target cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   comprising an anti-Fc a target cell .
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07-JUN-1995;
                                                                                                                                                                                              Sequence 1679
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               cagcttctggcttcaacattaaagactactatatgcactgggtgaagcagaggcctgaac
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DB; W73223.
cagcttctggcttcaacattaaagactcctatatgcactggttgaggcaggggcctgaac
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95US-0484172.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tumour cells - using a multi-specific molecule Fc receptor antibody and a portion which binds
                                                                                                                                 58.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Graziano
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                                                                                                              Score 425.2; per Pred. No. 7.3e-10; Mismatches
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                                                                                                                                 The invention relates to an antibody that binds a complex between prostate-specific antigen (PSA) and anti-chymotrypsin (ACT), and has an affinity for the complex which is at least 10 fold higher than the affinity for either PSA or ACT alone. The antibody is used in diagnostic assays to detect PSA-ACT in serum samples from patients. The levels of PSA-ACT complex increase in patients suffering from prostate cancer compared to the levels in patients with benign or no growths in the prostate. Therefore detection of PSA in complex with ACT is useful for the early detection of prostate tumours, by distinguishing between benign and malignant conditions of the prostate as well as for the management of patients with prostate cancer, such as the disclosure of metastasis and the monitoring of the PSA levels after treatment. The antibodies 'may' alone the monitoring of the PSA levels after treatment.
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also be used in immunotherapy, affinity chromatography and isolation purification of PSA-ACT. Unlike prior art antibodies which bind to PSA-ACT complexes but may also bind PSA or ACT alone, the present antibody is specific for PSA-ACT alone. Diagnostic assays using the antibodies are more accurate in diagnosing prostate cancer as they on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1548
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
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/cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US08_NEW_COMB.seq:*
                                                                                                                                                                                                                                                                              /cgn2_6/ptodata/2/pna/US6021_COMB.seq:*
/cgn2_6/ptodata/2/pna/US6022_COMB.seq:*
/cgn2_6/ptodata/2/pna/US6023_COMB.seq:*
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/cgn2_6/ptodata/2/pna/US6020_COMB.seq:*
cgn2_6/ptodata/2/pna/US60_NEW_COMB.seq:*
                                                                                                                                                                                                         COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

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; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(729)
US-09-297-181-1
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CURRENT FILING DATE: 1999-04-26
EARLIER APPLICATION NUMBER: PCT/FR97/01921
EARLIER FILING DATE: 1997-10-27
EARLIER APPLICATION NUMBER: FR96/13176
EARLIER FILING DATE: 1996-10-29
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
SEQ ID NO 1
LENGTH: 729
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Best Local Sim
Matches 729;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ANTI-P53 SINGLE-CHAIN ANTIBODY FRAGMENTS AND THEIR USES FILE REFERENCE: ST96030-US
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Pred. No. 1.8e-207;
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GENERAL INFORMATION
                                                                                                                                                                            Matches
                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (610) 454-3808 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NÚMBER: WO PCT/FI
FILING DATE: .29-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Fehlner Esq., Paul F.
REGISTRATION NUMBER: 35,135
REFERENCE/DOCKET NUMBER: ST9:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3839
                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 768 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 9
FILING DATE: 31-MAR-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: CONDITIONAL EXPRESSION SYSTEM NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SCHWEIG APPLICANT: SCHWEIG APPLICANT: TOCQUE,
 148
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                                                     STRANDEDNESS: dou
TOPOLOGY: linear
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                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 19426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Rhone-Poulenc Rorer Inc. STREET: 500 Arcola Road, Mailstop 3C43
                                                                                                                                   1 caggtgcagctgcagcagtctggggcagagcttgtgaggtcaggggcctcagtcaagttg
               gaaatcaaa 729
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35,135
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                                                                                                                                                                         Score 717.8; DB 13;
Pred. No. 4.2e-204;
Pred. No. 4.2e-7;
                                                                                                                                                                                                                                                                                                                                                                                                                  ST95021-US
                                                                                                                                                                                                        DB 13;
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APPLICANT: SCHOOLER, BRUND
APPLICANT: TOCQUE, BRUND
TITLE OF INVENTION: CONDITIONAL EXPRESSION S.
FILE REFERENCE: ST95021-US
CURRENT APPLICATION NUMBER: US/08/930,480A
CURRENT FILING DATE: 1998-01-21
PRIOR APPLICATION NUMBER: PCT/FR96/00477
PRIOR FILING DATE: 1996-03-29
PRIOR APPLICATION NUMBER: FR95/03841
PRIOR FILING DATE: 1995-03-31
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENCTH: 768
TYPE: DNA
ORGANISM: Artificial Sequence
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Query Match
Best Local Similarity
Matches 722; Conserv
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APPLICANT: BRACCO, Laurent APPLICANT: SCHWEIGHOFFER, Fabien
                                                                FEATURE:
OTHER INFORMATION: Description
OTHER INFORMATION: p53
OR-930-480A-4
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8 GAAATTAAA 756
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 Conservative
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              98.5%;
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              Score 717.8; DB 13
Pred. No. 4.2e-204;
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  Mismatches
                                                                                             Artificial
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RESULT 4
US-08-983-035-46
Sequence 46, Application US/08983035
Sequence 16, Application US/08983035
GENERAL INFORMATION:
APPLICANT: CONSEILLER, Emmanuel
APPLICANT: BRACCO, Laurent
ITILE OF INVENTION: P53 PROTEIN VAR
TITLE OF INVENTION: P53 PROTEIN VAR
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSEE; Rhone-Poulenc Rorer IN
ADDRESSEE; Rhone-Poulenc Rorer IN
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COUNTRY: USA5
ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatibl
OPERATING SYSTEM: PC-DOS/N
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                                                                  ADDRESSEE: Rhone-Poulenc
STREET: 500 Arcola Road,
CITY: Collegeville
STATE: PA
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; MOLECULE TYPE:
US-08-983-035-46
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Matches 7
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PROF APPLICATION DATA:
PRICATION NUMBER: WO FR96/01111
ETLING DATE: 17-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: Savitzky Esq., Martin F.
REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: ST95044-US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1NFORMATION:
TELEPHONE: (610) 454-3816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (610) 454-3808
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 749 base pairs
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APPLICATION NUMBER: FR 9
FILING DATE: 19-JUL-1995
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àgtggatcagggacagatttcacactgaaaatcaacagagtggaggctgaggatttggga 660
                                                         ttggatagtgatggaaagacatatttgaatttggttgttacagagggccaggccagtctcca<sup>*</sup> 540
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Pred. No. 1.
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Best Local S
Matches 721
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                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: WO FR96/01111
FILING DATE: 17-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: Savitzky.Esq., Martin F.
REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: ST95044-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (610) 454-3808 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 19-JUL-
PRIOR APPLICATION DATA:
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CLASSIFICATION:
61 teetgeacagettetggetteaacattaaagaetaetatatgeaetgggtgaagegg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 19426 4
                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
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                                                                                                                                                                                                                            NAME/KEY:
                                                                                                                                                                                                                                                                                                                          LENGTH:
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500 Arcola Road, Mailstop 3C43
                                                                                                                 Conservative
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98.9%;
                                                                                                              Score 716.2;
Pred. No. 1.8e
0; Mismatches
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Sequence 2, Application US/09203958
; Sequence 2, Application US/09203958
; GENERAL INFORMATION:
APPLICANT: Tibor Keler, et al.
TITLE OF INVENTION: CELLS EXPRESSING ANTI-FC RECFILE REFERENCE: MXI-099
; CURRENT APPLICATION NUMBER: US/09/203,958
; CURRENT FILING DATE: 1998-12-02
; EARLIER APPLICATION NUMBER: 60/067,232
; EARLIER FILING DATE: 1997-12-02
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2
; LENGTH: 1135
; TYPE: DNA
; TYPE: DNA
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Query Match
Best Local Similarity
Matches 608; Conserv
                                                                NAME/KEY: CDS
LOCATION: (74)..(1132)
-09-203-958-2
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              69.3%;
81.2%;
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 Score 505; DB 16;
Pred. No. 3.1e-140;
0; Mismatches 120;
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                         Length 1135;
  Indels
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SOFTWARE: Pat
SEQ ID NO 33
LENGTH: 819:
TYPE: DNA
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                                                                                             APPLICANT: KIKUCHI, YASUfumi
TITLE OF INVENTION: APOPTOSIS-INDUCING SINGLE-CHAIN
FILE REFERENCE: 055678/0102
CURRENT APPLICATION NUMBER: US/09/523,095A
CURRENT FILING DATE: 2000-03-10
                                                                                                                                                                            APPLICANT:
APPLICANT:
                                                               PRIOR EILING DATE: 1999-03-10
                                                                                                                                                   APPLICANT:
                                                        NUMBER OF SEQ ID
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                                                                                                                                                                                          FUKUSHIMA, Naoshi
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                                                                               2000-03-10
BER: JP 11-63557
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ORGANISM: FEATURE:

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Sequence 31, Application US/09523095A
; GENERAL INFORMATION:
; APPLICANT: FUKUSHIMA, Naoshi
; APPLICANT: UNO, Shinsuke
; APPLICANT: OH-EDA, Masayoshi
; APPLICANT: KIKUCHI, Yasufumi
; TITLE OF INVENTION: APOPTOSIS-INDUCIN
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NUMBER OF SEQ ID NOS: (
SOFTWARE: Patentin Ver
SEQ ID NO 31
LENGTH: 828
TYPE: DNA
ORGANISM: Mus sp.
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(822)
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Best Local Similarity 78.1%;
Matches 572; Conservative
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CURRENT APPLICATION NUMBER: US/0:
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: JP 11
PRIOR FILING DATE: 1999-03-10
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Pred. No. 3e-128;
0; Mismatches 157;
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US-09-297-181-3
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Best Local Similarity
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EARLIER FILING DATE: 1997-10-27
EARLIER APPLICATION NUMBER: FR96/13176
EARLIER FILING DATE: 1996-10-29
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/297,181
CURRENT FILING DATE: 1999-04-26
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                                              cagaaaccagggcagtctcctaaagtgctgatctactgggcatccactagggaatctgga
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6e-127;
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ORGANISM: Mus sp:
FEATURE:
NAME/KEY: CDS
LOCATION: (1)...(735)
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US-09-523-095A-39
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LENGTH: 741
TYPE: DNA
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Matches
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CURRENT APPLICATION NUMBER: US/09/523,095A
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: Jp 11-63557
PRIOR APPLICATION NUMBER: Jp 11-63557
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TITLE OF INVENTION: APOPTOSIS-INDUCING
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               ctcactttgtcggttaccattggacaaccagcctccatctcttgcaagtcaagtcagagc 477
                                                                                atggacctcagcagcctggcctctgaggactctgcggtctattactgtgcaagagggggt
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                                                             ggftcgggtggtggttcgggtggtggcggatcggatgttgtgatgacccaaagtcca
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OH-EDA, Masayoshi
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; LOCATION: (1)..(813)
US-09-523-095A-29
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US-09-523-095A-29
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GENERAL INFORMATION:
APPLICANT: FUKUSHIMA, Naoshi
APPLICANT: UNO, Shinsuke
APPLICANT: OH-EDA, Masayoshi
APPLICANT: KIKUCHI, Yasufumi
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SOFTWARE: PatentIn V
SEQ ID NO 29
LENGTH: 819
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Best Local Similarity
Matches 568; Conserv
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CURRENT APPLICATION NUMBER: US/09/523,095A
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: JP 11-63557
PRIOR FILING DATE: 1999-03-10
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Pred. No. 2.5e-126;
0; Mismatches 161;
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US-09-523-095A-25
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SEQ ID NO 25
LENGTH: 828
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: OH-EDA, MASAYUSHIA APPLICANT: KIKUCHI, YASUKUMI TITLE OF INVENTION: APOPTOSIS-INDUCING SINGLE-CHAIN FILE REFERENCE: 065678/0102
CURRENT APPLICATION NUMBER: US/09/523,095A
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: JP 11-63557
PRIOR FILING DATE: 199-03-10
PRIOR FILING DATE: 199-03-10
                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 568; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: FUKUSHIMA, Naoshi
APPLICANT: UNO, Shinsuke
APPLICANT: OH-EDA, Masayoshi
                                                                                                                                                                                                                                                                                     NAME/KEY: CDS
LOCATION: (1)..(822)
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GENERAL INFORMATION:
APPLICANT: Zhu, Zhenping
APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
ITILE OF INVENTION: Antibodies Specific to KDR and Use
FILE REFERENCE: Sequence Listings 1-34 for 381-71
CURRENT APPLICATION NUMBER: US/09/559,019
CURRENT FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
LENGTH: 714
TYPE: DNA
ORGANISM: Mouse
US-09-559-019-30
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                            tcctgcacaacttctggcttcaacattaaagacttctatatgcactgggtgaagcagagg
                                          tcctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaagcagagg
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77.3%;
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                                                                                                                                             Score 438; DB 21;
Pred. No. 3.1e-120;
0; Mismatches 145;
                                                                                                                                                                      Length 714;
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Sequence 29, Application US/09559019

Sequence 29, Application US/09559019

SEQUENCE INFORMATION:
APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
ITILE OF INVENTION: Antibodies Specific to KDR and Use
FILE REFERENCE: Sequence Listings 1-34 for 381-71

CURRENT APPLICATION NUMBER: US/09/559,019

CURRENT FILING DATE: 2000-04-25

NUMBER OF SEQ ID NOS: 34

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 29

LENGTH: 720

TYPE: DNA
ORGANISM: Mouse
US-09-559-019-29
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                                                                                              Query Match
Best Local Similarity
Matches 565; Conserv
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Pred. No. 9.3e-120;
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Query Match
Best Local Similarity
Matches 550; Conserv
                                                                                                      CURRENT APPLICATION NUMBER: PCT/US00/19843
CURRENT FILING DATE: 2000-07-21
PRIOR APPLICATION NUMBER: 60/145,512
PRIOR FILING DATE: 1999-07-23
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 16
LENGTH: 753
                                                                                                                                                                                                                                                                                       Sequence 16, Application PC/TUS0019843 GENERAL INFORMATION:
                                                                                                                                                                                                                        APPLICANT: Deikman, Alan B.
TITLE OF INVENTION: Recombinant Antibody Directed Against Human Sperm
TITLE OF INVENTION: Antigen
                                                                                                                                                                                                                                                 APPLICANT: Herr, John C.
APPLICANT: Norton, Elizabeth J.
APPLICANT: Deikman, Alan B.
                                                                                                                                                                                                             FILE REFERENCE:
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Search completed: February 18, 2001, 01:29:16 Job time: 18114 sec

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Result
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2: /cgn2_6/ptodata/2

3: /cgn2_6/ptodata/2

4: /cgn2_6/ptodata/2

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US-08-902-486-10

US-08-97-689-17

US-08-752-844-5

US-08-752-844-5

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US-08-465-4738-10

US-08-465-4738-10

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	Sequence 71, Appl	Sequence 8, Appli				Sequence 18, Appl		Sequence 23, Appl	Sequence 9, Appli	Sequence 8, Appli	Sequence 6, Appli	Sequence 13, Appl	Sequence 6, Appli	•	Sequence 11, Appl	Sequence 27, Appl	Sequence 9, Appli

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; FEATURE: mat_peptide; NAME/KEY: mat_peptide; LOCATION: (95)...(895); OTHER INFORMATION: Coding sequence for genetically engineered single; OTHER INFORMATION: chain antibody - 5109 VH - VL. US-09-184-658-9
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US-09-184-658-9
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APPLICANT: Otterness, Ivan G.
APPLICANT: Mezes, Peter S.
APPLICANT: Downs, James T.
APPLICANT: Johnson, Kimberly S.
APPLICANT: Johnson, Kimberly S.
TITLE OF INVENTION: Biological Media
FILE REFERENCE: PC9946-A
                                                                                                                                                                                                                                                                                                      SEQ ID NO 9
LENGTH: 907
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/184,658
CURRENT FILING DATE: 1998-11-02
EARLIER APPLICATION NUMBER: 60/065,423
EARLIER FILING DATE: 1997-11-13
NUMBER OF SEQ ID NOS: 69
SOFTWARE: Patentin Ver. 2.0
                                                                                                                              NAME/KEY: sig_peptide
LOCATION: (29)..(94)
OTHER INFORMATION: Engineered
OTHER INFORMATION: methionine
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                                                                                                                                                                                                                                            Description of Artificial Sequence: single chain antibody.
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is coded for most
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likely by
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2 aggtgcagctgcagcagtctggggcagagcttgtgaggtcaggggcctcagtcaagttgt

Conservative

59.7%; 75.6%;

Score 435; DB 3; Pred. No. 5e-119; Mismatches 175; DB 3;

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APPLICATION NUMBER: US/08/652,507
ETLING DATE: 02-Jul-1996
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: ATTHUR R. Crawford
REGISTRATION NUMBER: 25,327
REFERENCE/DOCKET NUMBER: 117-211
                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                             TITLE OF INVENTION: ANT
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon & Va
                                                                                          ZIP: 22201-4714

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                         CITY: Arlington
STATE: Virginia
COUNTRY: USA
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 810 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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Best Local S
Matches 568
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TELEPHONE: 703-816-4000
TELEFAX: 703-816-4100
                                            <sup>₩</sup> 721
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                                                         GGCACTTCTCCCAAACTCTGGATTTATAGCACATCCAACCTGGCTTCTGGAGTCCCTGCT
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76.78;
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Pred. No. 3.2e-118;
0; Mismatches 143;
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RESULT 3 US-08-661-052-15 ; Sequence 15, Application

US/08661052

Patent No.

GENERAL INFORMATION:

Yashwant M. Deo Joel Goldstein Robert Graziano

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; LOCATION:
US-08-661-052-15
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Best Local Similarity 76.4
Matches 561; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCE CORRESPONDENCE ADDRESS:

CORRESPONDENCE LAHIVE & COCKFIELD

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LENGTH: 1679 base pairs
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REFERENCE/DOCKET NUMBER: MX
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617,722,7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 07-JUNE-199
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 07-JUNE-1995
                                                                                                                     1086 TGAGCAGCCTGACATCTGAGGACACTGCCGTCTATTATTGTAATGAGGGGACTCCGACTG 1145
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296 acggggatgctttgggactattggggccaagggaccacggtcaccgtctcctcaggtggag
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Pred. No. 5e-116;
0; Mismatches 143;
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US-08-875-811-52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: _Rybak, Susanna
        INFORMATION FOR SEQ ID NO:
                                                        FILING DATE: 21-FEB-1996
ATTORNEY/AGENT (INFORMATION:
NAME: FAIIS, SUSAN K.
REGISTRATION NUMBER: 41,739
REFERENCE/DOCKET NUMBER: 015;
TELECOMMUNICATION INFORMATION:
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APPLICANT: N
APPLICANT: B
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                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1428
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                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
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PPLICANT: Wlodawer, Alexander
ITLE OF INVENTION: Recombinant Ribonuclease Proteins
                                                                                                                                                                                                                       APPLICATION NUMBER: WO P
FILING DATE: 19-FEB-1997
                                                                                                                                                                                                                                                                                      FILING DATE: · 19-FEB-1998 CLASSIFICATION: 435
                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
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                         (415)
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                                           (415) 576-0200
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Dianne L.
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RESULT 5
US-08-875-811-54
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; Patent, No. 6045793
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LOCATION: 1..1095;
OTHER INFORMATION:
US-08-875-811-52
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Best Local :
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STRANDEDNESS: single
TOPOLOGY: linear
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Pred. No. 7.3e-111;
0; Mismatches 201;
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LOCATION: 1..1098
OTHER INFORMATION:
US-08-875-811-54
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                           Matches 528;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/011,
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Faris, Susan K.
REGISTRATION NUMBER: 41,739
REFERENCE/DOCKET NUMBER: 0152:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 'SEQUENCE CHARACTERISTICS:
LENGTH: 1098 base pairs
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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APPLICANT: Newton, Dianne
APPLICANT: Boque, Lluis
 244
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                             241 ctgcagctcagcatggcatctgaggacactgccgtctattattgtaatttttacggg 300
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STRANDEDNESS: sing
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CLASSIFICATION:
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                                                                                               gccccgaagtt.ccagggcaaggccactatgactgcagacacatcctccaatacagcctac 240
                                                                                                                                  CCAGGAAAGGGTTTAAAGTGGATGGGCTGGATAAACACCTACACTGGAGAGTCAACATAT
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                                                                    GCTGATGACTTCAAGGGACGGTTTGCCTTTTCTCTAGAAACCTCTGCCAGCGCTGCCTAT
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Two Embarcadero Center, F
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21-FEB-1996
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Pred. No. 7.3e-111;
0; Mismatches 201;
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Eighth Floor
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US-08-331-398A-33
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MEDIUM TYPE: Floppy utt...
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-POS/MS-DOS
OPERATING FOR PATENTIAL Release #1.0,
TOTTWARE: PATENTIAL TOP /331,39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 33, Application US/08331398A Patent No. 5608039
                                            APPLICATION NUMBER: US/08/331,398A FILING DATE: 28-OCT-1994 CLASSIFICATION: 435
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 07/767,331
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US 07/596,289
PILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                      NUMBER OF SHAPESS:
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew
ADDRESSE: Townsend Plaza, Steuart Street Plaza
                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David
APPLICANT: Brinkmann, Ulrich
APPLICANT: Pai, Lee
TITLE OF INVENTION: Single Chain B3 Antibody Fusion Proteins
TITLE OF INVENTION: and Their Uses (as amended)
NUMBER OF SEQUENCES: 68
NUMBER OF SEQUENCES: 68
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STREET: U...
STREET: On...
STRY: San F
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    REFERENCE/DOCKET NUMBER:
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Best Local Similarity
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TYPE: nucleic acid
STRANDEDNESS: single
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Pred. No. 1.7e-97;
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US-08-331-397B-33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTY 94105-1492
ZIP: 94105-1492
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACTIBLE
COMPUTER: TOTEM: PC-DOS/MS-DOS
COMPUTER: TOTEM: PC-DOS/MS-DOS
COMPUTER: TOTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-0CT-1990
ATTORNEY/AGENT INFORMATION:
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APPLICANT: Benhar, Ital
APPLICANT: Chi
                                                                                                                            FEATURE:
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CORRESPONDENCE ADDRESS:
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TOPOLOGY: line
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 0 FILING DATE: 30-SEP-1991
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CITY: San Francisco
                                                                                                                                                                                                                                NAME/KEY:
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protein of B3 monoclonal antibody
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                 Score 361.8; DB 2; Pred. No. 1.7e-97;
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    Mismatches
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                                                                                       GENERAL INFORMATION:
APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David J
APPLICANT: Brirkmann, Ulrich
APPLICANT: Pal, Lee
                                                                                                                                                                                         Sequence 33, Application US/08759804A Patent No. 5990296
                                            TITLE OF INVENTION: Tumor-S
TITLE OF INVENTION: Fusion
NUMBER OF SEQUENCES: 68
             CORRESPONDENCE ADDRESS:

ADDRESSEE: 'Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                     713 ccaagctggaaatcaaa
| ||||||||||||
| 751 CAAAGCTGGA&ATTAAA
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Two Embarcadero Center, Eighth
                                                             Tumor-Specific Antibody Fragments Fusion Proteins, and Uses Thereof
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COUNTRY: STATE:

San California

Francisco

Y: USA 94111-3634

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US-08-759-804A-33
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FILING DATE: 28-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
NAME: Weber: Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 015280-126140US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200 TELEPAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 33:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 772 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: 1..772
OTHER INFORMATION:
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   331
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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LOCATION:
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                                                                      cctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaagcagaggc 121
                       tttacggggatgctttggactattggggccaagggaccacggtcaccgtctcctcaggtg
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                                                                                                                                                CAGACACTGTAAAGGGCCGGTTCACCATCTCCAGAGACAATGCCAGGAACACCCTCTACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "Single-chain antibody fusion protein of B3 monoclonal antibody Variable Heavy chain (V-H) and Variable Light chain (V-L) Fv region joined by a (Gly-4Ser)-3 peptide linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 361.8;
Pred. No. 1.7
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                                                                                                                                            REFERENCE/DOCKET NUMBER: SC
TELECOMMUNICATION: INFORMATION:
TELEPHONE: 703-243-6333
TELEPHX: 703-243-6410
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/800,198
FILING DATE: 13-FEB-1997
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
ARME: HAMLET FRING, DIANA
REGISTRATION NUMBER: 33,302
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 720 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: WELS, WINFRIED S.
APPLICANT: SCHMIDT, MATHIAS
APPLICANT: VAKALOPOULOU, EVANGELIA
                 MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: SCHNEIDER, DOUGLAS
TITLE OF INVENTION: GROWTH FACTOR RECEPTOR
NUMBER OF SEQUENCES: 17
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ZIP: 222
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                      TOPOLOGY:
                                                                    TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 atttgggagtttattattgctggcaaggtacacattctccgctcacgttcggtgctggca 712
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Sequence 7, Application US/09296595A
Patent No. 6129915
GENERAL INFORMATION:
APPLICANT: WELS, WINFRIED S.
APPLICANT: SCHMIDT, WATHIAS
APPLICANT: VARALOPOULOU, EVANGELIA
APPLICANT: SCHMIDER, DOUGLAS
APPLICANT: SCHMIDER, DOUGLAS
TITLE OF INVENTION: GROWTH FACTOR RECEPTOR AN
FILE REFERENCE: SCH-1576 D.
FILE REFERENCE: SCH-1576 D.
CURRENT APPLICATION NUMBER: US/09/296,595A
CURRENT FILING DATE: 1999-04-23
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Mismatches 196;
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RESULT 11
US-08-463-163-2
; Sequence 2, Application U
; Patent No. 5696237
; GENERAL INFORMATION:
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LOCATION: 1.1797; OTHER INFORMATION: US-08-463-163-2
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Matches 502; Conserv
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
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NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
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181 gccccgaagttccagggcaaggccactatgactgcagacacatcctccaatacagcctac 240
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OPERATING SYSTEM:
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WIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 05-JUN
                                                                                                        64
                                                                                                                                     61 tcctgcacagcttctggcttcaacattaaagactactatatgcactgggtgaagcagagg 120
                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 06/227,227 FILING DATE: 22-JAN-1981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
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One Market Plaza, Steuart Street Tower
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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24-SEP-1986
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05-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 31,
                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                 APPLICATION NUMBER: US/0 FILING DATE: '28-OCT-1994 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Single Character of INVENTION: and Their
 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        244 ATGCAACTGAGCAGCCTGACATTTGAGGACTCTGCAGTCTATTACTGTGCAAGAGGGGGG
                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                           CITY: San Francisco
STATE: California
                  APPLICATION NUMBER: US 07/767,331 FILING DATE: 30-SEP-1991
                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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One Market Plaza, Steuart Street
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Willingham, Mark
FitzGerald, David
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; OTHER INFORMATION:
US-08-331-398A-31
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Hunter Tom
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alliggagittattattgctggcaaggtacacattctccgctcacgttcggtgctggca
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                                                                      tcactggcagtggatcagggacagatttcacactgaaaatcaacagagtggaggctgagg 652
                                                                                                                                                           agtotocaaagogootaatotatotggtgtotaaactggaototggagtocotgacaggt 592
                                                                                                                                                                                                                                                                    CTCCATTGAGTTTACCTGTCACCCCGGGAGAGCCGGCCTCCATCTCTTGCAGATCTAGTC
                                                                                                                                                                                                                                                                                                                                          GAGGCGGATCCGGTGGTGGCGGATCTGGAGGTGGCGGAAGCGATGTGCTGATGACCCAGT 421
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68.28;
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Pred. No. 9.9e-94;
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                                                                                                                                                                                                                                                                                                           Best
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILLING DATE: 12-OCT-1990
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US.
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MEDITM TYPE: Floppy disk
COMPUTER: IBL PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
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LOCATION: 1:.738
OTHER INFORMATION:
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STRANDEDNESS: sing
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                                                                                                                                                                                                                                                                                                           Local Similarity
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   CCTGTGCAACCTCTGGATTCACTTTCAGTGACTATTACATGTATTGGGTTCGCCAGGCCC
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One Market Plaza, Steuart Street
                                                                                                                                                                                                                                                                    Conservative
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Specific Antibody Fragments, Fusion Proteins, and
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9.9e-94;
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RESULT 14
US-08-759-804A-31
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APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David
APPLICANT: Brinkmann, Ulrich
APPLICANT: Pai, Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 31, Patent No. 5
                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                 TITLE OF INVENTION: TUMOR'S
TITLE OF INVENTION: Fusion
NUMBER OF SEQUENCES: 68
CORRESPONDENCE ADDRESS:
                                                                                                                                                                        STATE: C
 APPLICATION NUMBER: FILING DATE: 03-DEC
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                                                                                                                                                                                       San Francisco
California
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                                                                                                                                                                                                                                                                                                                                            Brinkmann, Ulrich
Pai, Lee
   03-DEC-1996
                                                                                                                                                                                                                                                                                                                      Tumor-Specific Antibody Fragments
                     US/08/759,804A
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 738 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 503; Conserv
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 30-SEP-15
PRIOR APPLICATION NUMBER: US
APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 12-OCT-1990 ATTORNEY/AGENT INFORMATION:
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OTHER INFORMATION:
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agtotocaaagogocotaatotatotggtgtotaaactggactotggagtocotgacaggt
                                                                               agagcctcttggatagtgatggaaagacatatttgaattggattgttacagaggccaggcc
                                                                                                                                  CTCCATTGAGTTTACCTGTCACCCCGGGAGAGCCGGCCTCCATCTCTTGCAGATCTAGTC
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Score 349; DB Pred. No. 9.9e 0; Mismatches

DB 2; L).9e-94; nes 225;

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US-08-279-772A-7
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                                                                                                            Query Match 47. Best Local Similarity 68. Matches 501; Conservative
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                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27,386
REGISTRATION NUMBER: 11-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-251-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                            TELEFAX: 608-251-9166 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Methor
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
67
                             61
                                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: United ZIP: 53701-2113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Madison
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 TCCTGCAAGGCTTCTGGCTACACCTTTACTAGCTACAGGATGCACTGGGTAAAACAGAGG
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/ENTION: Method for Producing Antibodies in Plant
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                                                                                                                                                                                                                                        DNA (genomic)
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                                                                                                           Score 347.6; DB 3;
Pred. No. 2.5e-93;
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GAGCTCAA 716
                                             ACTTATTACTGCCATCAAAGGAGTACTTACCCACTCACGTTCGGGTTCTGGGACCAAGCTG
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Search completed: February 17, 2001, 22:59:21 Job time: 35112 sec

1

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 12, 2001, 15:22:50; Search time 26.51 Seconds (without alignments)
622.401 Million cell updates/sec

Title: US-09-297-181-2
Perfect score: 1289
Sequence: 1289
Sequence: 1289
Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5
Searched: 195891 seqs, 67900655 residues
Total number of hits satisfying chosen parameters: 195891
Minimum DB seq length: 0
Maximum DB seq length: 2000000000
```

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

PIR_66:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

SUMMARIES

A; Reference number: JC5322; MUID: 97168950

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447	449	450	451	453	458.5	460	460.5	461	462	463	464.5	465	468.5	469	469
34.7	34.8	34.9	35.0	35.1	35.6	35.7	35.7	35.8	35.8	35.9	36.0	36.1	36.3	36.4	36.4
112	132	115	131	122	114	112	114	133	114	118	144	103	140	133	91
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S38719	S40322	S38715	в39276	S40338	S49572	A31807	B49002	A24452	A47271	S40374	в30502	PH1056	S22658	K2HURP	S42186
					i										
Ig light chain V r	Ig kappa chain – h	Ig kappa chain V r	Ig light chain pre	chain	Ig kappa chain pre	chain		Ig kappa chain pre	nitrophenyl phosph	Ig kappa chain - h	Ig heavy chain V r	Ig light chain V r	Ig kappa chain pre	Ig kappa chain pre	Ig kappa chain V r

ALIGNMENTS

RESULT S41374

	<pre>single chain Fv antibody - mouse C:Species: Mus musculus (house mouse)</pre>	
	<pre>C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 06-Jan-1995 C:Accession: S41374 \ C:Acces</pre>	
	R; Artsaenko, O.; Weiler, E.W.; Muentz, K.; Conrad, U.	
	Submitted to the EMBL Data Library, January 1994 A;Description: Construction and functional characterization of a single chain Fv anti	
	orelim:	
	A;Molecule type: DNA A;Residues: 1-249 <art> A:Cross-references: EVMI: 224480</art>	
	Winters teterances Tube: 753400	
	Query Match 70.8%; Score 912.5; DB 2; Length 249;	
	nes 179; Conservative 25; Mismat	
	QY 1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60	
	Db 1 QVQLQQSGAELVRPGASVKLSCTASGFNFKDDYIHWVKQRPEKGLEWIARIAPASGNVKY 60	
	Qy 61 APKFQGKATMTADISSNTAYLQLSSLASEDTAVYYCNFYGDALDYWGQGTTVTVSS 116	
	Db 61 VPREQDKATITADTSSNTAYLLLSSLTSEDTAVYYCARRDTLY-TSLGYWGQGSTVTVSS 119	
	QY 117 GGGGSGGGSJGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRP 176	
	Db 120 RGGGSGGGGSGGGSDIELTQSPPSVVVIPGESVSISCRSSKSLLYSDGDSYLFWFLQRP 179	
	QY 177 GQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGA 236	
	Db 180. GOSPOLLIYRMSNLASGVPDRFSGSGSGTSFTLRISRVEAEDVGVYYCMQHREXPLTFGA 239	
	QY 237 GTKLEIK 243	
	Db 240 GTKLELK 246	
• . •	RESULT 2 · JC5322	
	p53 specific single-chain antibody Pab421 - human	
,	C;Date: 15-May-1997 #sequence_revision 15-May-1997 #text_change 18-Jul-1997	
	R:Jannot, C.B.: Hynes, N.E.	
	Biophys. Res. Commun. 230, 242-246, 1997	
	A; Title: Characterization of scFv-421, a single-chain antibody targeted to pos-	

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C;Date: 19-Jan-1990 "Value C;Date: 19-Jan-1990 "Value C;Accession: A56446
R;Tang, P.M.; Foltz, L.A.; Mahoney, W.C.; Schueler, J. Biol. Chem. 270, 7829-7835, 1995
A;Title: A high affinity digoxin-binding protein dir A;Reference number: A56446; MUID:95229583
A;Accession: A56446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Accession: JC5322
A; Accession: Lype: max,
A; Mclecule type: Max,
A; Residues: 1-233 <JJ
A; Experimental source
C; Comment: This prot
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A; Residues: 1-268 < TAN>
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g heavy chain V region (3H-3H scFv) - mouse (strain BALB/C)
Species: Mus musculus (house mouse)
Date: 19-Jan-1996 #sequence_revision 19-Jan-1996 #text_cha
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Experimental source:
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;Residues: 1-233 <JAN>
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Best Local
237
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                                KLEIK 243
                                                                                                                                                                     GGSGGGGSGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRPGQ 178
                                                                                                                                                                                                        DPKFQGKATIAADTSSNTAYLQLSSLTSEDTAVYYCASYYLTRYENYWGQGTTVTVSSGG
                                                                                                                                                                                                                           APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC-NFYGDALD-YWGQGTTVTVSSGG 118
                                                                                                                                                                                                                                                                                              QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
KLEIK
                                                                  SPKLWYYTSHLPPGVPARFSGSGSGNSYSLTISSMEGEDAATYYCQQFTSSPFTFGSGT
                                                                                                 SPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGT 238
                                                                                                                                     GGSGGGDSGGGGSDIELTQSPAIMSASLGEKVTMSCRASSSV-----NFIYWYQQKSDA 176
                                                                                                                                                                                                                                                                          QVKLQESGAELVKPGASVKLSCTTSGFNIKDTYMHWVKQRPEQGLEWIGRIAPANGITKY 62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QGKATMTADTSSNTAYLQLSSLASEDTAVYYCNFYGDALDYWGQGTTVTVSSGGGGSGGG
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54; Conservative
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Pred. No. 3.2e-51;
0; Mismatches 49
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Pred. No. 1.
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Mismatches 44
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RESULT C32513

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response:

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Ig heavy chain precursor v region (MRL-histone 7H) C; Species: Mus musculus (house mouse) C; Date: 04-Dec-1992 *sequence_revision 04-Dec-1992 *C; Accession: S04576
                                                                                                                            RESULT
S04576
                                                                                                                                                                                                    В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ig kappa chain precursor V region (BXW14) - mouse C. Species: Mus musculus (house mouse) C. Date: 21-May-1990 #sequence_revision 31-Dec-1991 C; Accession: C32513 . R. Kofler, R.; Strohal, R.; Balderas, R.S.; Johnson J. Clin. Invest. 82, 852-860, 1988 A; Title: Immunoglobulin kappa 119ht chain variable A; Reference number: A94689; MUID:88331394
   Eur. J. Immunol. 1
A;Title: Molecular
                    R; Kofler, R.; Noonan, Eur. J. Immunol. 17,
                                                                                                                                                                                                                                   QΥ
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-95/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-113 <MAT>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Reference number: A; Accession: F30560
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A; Residues: 1-132 KKOF>
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Best Local Similarity
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Best Local (
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an, D.J.; Strohal, R.; 7, 91-95, 1987 analysis of the murino
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94.6%;
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Pred. No. 5
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Pred. No. 2.1e-33;
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 lupus-associated
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.5e-33;
                                        R.S.;
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     anti-self
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                                          F.J.;
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C;Accession: A36259
R;Zilber, B.; Scherf, T.; Levitt, M.; Anglister, J.
Biochemistry 29, 10032-10041, 1990
A;Title: NMR-derived model for a peptide-antibody complex.
A;Reference number: A36259; MUID:91104915
                                                                                                                                                                                       ig kappa chain V region (TE34) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 18-Jan-1991 #sequence_revision 18-Jan-1991 #text_change 21-Jan-2000
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A; Cross-references: GB:L34775
A; Note: authors translated the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Title: Molecular cloning of a proteolytic antibody light chain A;Reference number: A55491; MUID:95096089
A;Accession: A55491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            proteolytic antibody light chain - mouse
C;Species: Mus musculus (house mouse)
C;Date: 03-Mar-1995 #sequence_revision 03-Mar-1995 #text_change 21-Jan-2000
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A;Accession: S04576
                    A; Molecule type: mRNA
A; Residues: 1-112 <ZIL>
                                                                         A; Reference number: A; Accession: A36259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Gao, Q.S.; Sun,
J. Biol. Chem. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A55491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
                                                      A;Status: preliminary
                                                                                                                                                                                                                                                    A36259
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Cross-references: GB:M30458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Accession: A55491
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        104;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     un, M.; Tyutyulkova, S.
269, 32389-32393, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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88.9%;
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92.9%;
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lin V region; immunoglobulin homology
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GB:M30459; GB:M30480; GB:M30481;
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Pred. No. 2.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 541; DB 2;
Pred. No. 3.9e-32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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2.3e-32;
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GB:M30482; GB:M30483
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C:Date: 201709 ;
(C:Date: 2017
C;Superfamily: immunoglobulin V region; im C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <I
                                                                                                                                                                                                                                                      A; Title: Reshaping a human r
A; Reference number: 515672;
A; Accession: 515672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig heavy chain V region - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
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C;Keywords: heterotetramer; immunoglobulin
F;16-95/Domain: immunoglobulin homology <II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: $20706
A; Accession: $20709
A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig kappa chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 20-Feb-1995 #sequence_revision
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                                                                                                               A; Cross-references: EMBL: 8X58835; NID: g51978; PIDN: CAA41644.1; PID: g51979
                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-116 <TEM>
                                                                                                                                                                                                                           A; Status: preliminary; nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                         Bio/Technology 9, 265-271,
                                                                                                                                                                                                                                                                                                                                                                                                               R; Tempest, P.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Accession:
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A; Residues: 1-111 <BRE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   102;
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                                                                                                                                                                                                                                                                                                                                                                                                        Brenner, P.; Lambert, M.; Taylor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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91.9%;
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92.9%;
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Pred. No. 5.4e-32;
2; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 537; DB 2; Pred. No. 7.5e-32;
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   <IMM>
                                                                         immunoglobulin
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                                                                                                                                                                                                                               not
                                                                                                                                                                                                                               shown
                                                                                                                                                                                                                                                                                                                                                                                                        G.; Furze, J.M.; Carr, F.J.;
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Query Match Best Local Similarity

41.0%; 84.3%;

Score Pred.

528; DB 2; No. 3.4e-31;

Length 116

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C;Accession: $52445
R;Berdoz, J.; Kraehenbuhl, J.P.
submitted to the EMBL Data Library, November 1994
A;Description: Specific amplification by the polyn
A;Reference number: $52445
A;Accession: $52445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              F;55-61/Region: complementarity-determining F;62-93/Region: framework 3
F;94-102/Region: framework 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              F;16-95/Domain: immunoglobulin homology <IM
F;24-39/Region: complementarity-determining
F;40-54/Region: framework 2
F;55-61/Domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Roth J. Exp. Med. 171, 265-297, 1990 A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and A;Reference number: PLO231; MUID:90111618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ig kappa chain V region (anti-DNA, D23VK) - mouse (fragment)
C;Speciles: Mus musculus (house mouse)
C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
C;Accession: PL0273
                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-137 <BER>
                                                                                                                                                                                                                                                                                                                                                                                 Ig heavy chain V region precursor - mouse
C:Species: Mus musculus (house mouse)
C:Date: 08-May-1995 *sequence_revision 21-Jul-1995 *text_change 21-Jan-2000
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A; Residues: 1-112 <SHL>
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                                                                                                                                                                                  A;Cross-references: EMBL:X82690; NID:g673439; PIDN:CAA58011.1; PID:g673440
                                                                                                                                                                                                                                          A; Status: preliminary
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                                                                                                    ;Superfamily: immunoglobul;
Keywords: heterotetramer;
Query Match
Best Local S
                                                                           34 (115/Domain:
                                                                                                                                                               Genetics:
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Best Local Similarity
Matches 103; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRPGQSPKRLIYLVSKLD 191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SGVPDRFTGSGSGTDFTLKISRVEAEDLGVYYCVQGTHFPWTFGGGTKLEIK 112
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  Similarity
                                                                                                  immunoglobulin V region; imcerotetramer; immunoglobulin
                                                                             immunoglobulin
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92.08;
  40.1%;
83.1%;
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Pred. No. 3.9e-31;
    Score
Pred.
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No. 2.
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  DB 2;
.5e-30;
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                    Length 137;
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A; Molecule type: mRNA.
A; Residues: 7-120 <ROC1>
A; Residues: 7-120 <ROC1>
A; Residues: 7-120 <ROC1>
A; Roross-references: EMBL: X01820; NID: g51833; PIDN: CAA25962.1; PID: g1333983
A; Note: this sequence was determined from the differentiated gene
:R; Rocca-Serra, J.; Mazie, J.C.; Moinier, D.; Leclercq, L.; Somme, G.; Theze,
J. Immunol. 129, 2554-2558, 1982
A; Title: The limited diversity of the mouse gamma-chains anti-GAT repertoire
A; Reference number: $07453; MUID: 83058021
A; A; Accession: $07453
A; Molecule type: protein
A; Residues: 1-43 <ROC2>
A; Residues: 1-43 <ROC2>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: GB:M25996; NID:g197109; PIDN:AAA38911.1; PID:g197110 A;Note: the authors translated the codon CGC for residue 51 as Leu, and C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;16-95/Domain: immunoglobulin homology <IMM>
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A;Title: Early rearrangements of genes encoding murine immunoglobulin kappa-chains, A; Reference number: A33730; MUID:89367325
A; Accession: A33730
                                                                                                                                                                                                                                                                                                                                                             A:Title: Analysis of antibody diversity: V-D-J mRNA nucleotide hypervariable regions. A:Reference number: S03471; MUID:84057768 A:Accession: S03471
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EMBO J. 2, 867-872,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Accession: S03471; S07453
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C;Species: Mus' musculus (house mouse)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 20-Jun-2000
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A;Molecule type: DNA
A;Residues: 1-101 <LAW>
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C; Species: Mus musculus (house mouse)
C; Date: 09-Mar-1990 #sequence_revision 18-Sep-1992
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97; Conservat
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Pred. No. 4.8e-30
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A;Description: Immunoglobulin variable heavy and light chain cDNA sequences for two anti A;Reference number: S31577
A;Accession: S31577
A;Accession: S31577
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C;Species: Mus musculus (house mouse)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                          ,;Status: preliminary
,Molecule type: mRNA
,Residues: 1-131 <REC>
,Cross-references: EMBL:Z19575; NID:g53983; PIDN:CAA79627.1; PID:g53984
;Cross-references: EMBL:Z19575; NID:g53983; PIDN:CAA79627.1; PID:g53984
;Superfamily: immunoglobulin V region; immunoglobulin homology
;Keywords: heterotetramer; immunoglobulin
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Matches 97; Conservative
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MEDLINE=73166638;
                                                                                   Putnam F.W., Whitley E.J. Jr., I
"Amino acid sequence of a kappa
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COMPLEMENTARITY-DETERMINING
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P01631;

21-JUL-1986 (Rel. 01, Created)

T 21-JUL-1986 (Rel. 01, Last sequents)

(Rel. 38, Last annous)

TV V-II REGION 26-1
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Glenner G.G.
"Structural
                                                                                                                                                                                                                         MEDLINE-831/022, ... MAIGOILES M.N.; ROVOTHY J., MAIGOILES M.N.; RAMINO acid sequence of the light canti-digoxin hybridoma antibody."; Biochemistry 22:1153-1158(1983).
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-!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE INV (1)
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                                                                                         DOMAIN
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                FRAMEWORK 3.
COMPLEMENTARITY-DETERMINING
FRAMEWORK 4.
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COMPLEMENTARITY-DETERMINING
FRAMEWORK 2.

COMPLEMENTARITY-DETERMINING
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COMPLEMENTARITY-DETERMINING
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01-JAN-1988 (Rel. 06, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IG KAPPA CHAIN V-II REGION GM607 PRECURSOR (FRAGMENT).
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                                                                                                                                                                                                                                                                                                                                                    the European Bioinformatics Institute. The use by non-profit institutions as love modified and this statement is not removed. entitles requires a license agreement (See or send an email to license@lsb-sib.ch).
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01-JAN-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=84191506; PubMed=6325927;
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SKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGTKLEIK 243
                      GSSGDIVMTQSPLSLPVTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPQQSPQLLIYLG
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Pred. No. 6.7e-26;
5; Mismatches 20
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Pred. No. 2.7e-26;
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                                                                                                                                                                COMPLEMENTARITY-DETERMINING 3. FRAMEWORK 4.
                                                                                                                                                                                                  COMPLEMENTARITY-DETERMINING 2.
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                                     RESULT 7
KV2A_HUMAN
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Best Local
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            KV2A_HUMAN
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IG HEAVY CHAIN V REGION B1-8/186-2 PRECURSOR.
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P01751; P01752; +
21-JUL-1986 (Rel. 01, Created)
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MEDLINE-81234548;
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
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                                                                                                    61 APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC---NFYGDA-LDYWGQGTTVTVSS
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                                                                                      NEKFKSKATLIVDKPSSTAYMQLSSLTSEDSAVYYCARYDYYGSSYFDYWGQGTTLTVSS
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Pred. No. 8.2e-26;
5; Mismatches 20
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          INTERPRO; IPR003006; -.
PFAM; PF00047; 1g; 1.
Immunoglobulin V region; Antiarsonate antibody; Hybridoma.
NON_TER 120 120
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                                                                                                                                                                                                                                    Mus musculus (Mouse)
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                              21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last ser
15-JUL-1999 (Rel. 38, Last and
IG HEAVY CHAIN V REGION 36-65
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Immunoglobulin V region; Bence-Jones proter
24 95 BY SIMILARITY.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUL-1986 (Rel. 01, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) IG KAPPA CHAIN V-II REGION CUM.
                                                                                                                                                               Marshak-Rothstein A.;
                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                           Mammalia;
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Naturwissenschaften 56:195-20
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                                                                                                                         he genetic basis of antibody production: liotype response of the strain A mouse."; Inmunol. 12:1023-1032(1982).
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                                                                                   MISCELLANEOUS: FROM ANALYSIS OF THE SIZES OF SEVERAL OTHER DIFFERENTIATED GENES THAT HYBRIDIZE TO THIS ONE, THE AUTHOLOCONCLUDE THAT ALL OF THESE V REGIONS HAVE REARRANGED TO TH
                                                           SEGMENT, JH2.
A02028; HVMSG7.
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RPRO; IPR003006; -.
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115 AA;
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12676 MW;
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 13307 MW;
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205(1969).
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Pred. No. 1.5e-25;
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 FF04E4A167B654AF CRC64;
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P01755;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
16 HEAVY CHAIN Y REGION $43 PRECURSOR.
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CHAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                      MISCELLANEOUS: THE GAMMA-2A CHAIN MENA WAS CLONED FROM A HYBRIDOMA MAKING ANTIBODIES TO THE HAPTEN (4-HYDROXY-3-NITROPHENYL)ACETYL
  QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
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01, Last sequence update)
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                                    31.0%;
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                          Score 399; DB 1;
Pred. No. 5.3e-24;
0; Mismatches 28
                                                                                                             COMPLEMENTARITY-DETERMINING FRAMEWORK 3.
D SEGMENT.
JH2 SEGMENT.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          contribution to the evident in a gamma
                                                                                                                                                                           FRAMEWORK
                                                                                                                                                                                      COMPLEMENTARITY - DETERMINING 1.
                                                                                                                                                                                                    FRAMEWORK
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Matches 76
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                                      Homo sapiens (Human)
Eukaryota; Metazoa;
                                                                                                          21-JUL-1986 (Rel.
21-JUL-1986 (Rel.
15-JUL-1999 (Rel.
                                                                                                                                                                                   KV2C_HUMAN
P01616;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   between the Swiss Institute of Bioinf
the European Bioinformatics Institute
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Science 216:309-311(1982).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=82152818; PubMed=6801765; Sims J., Rabbitts T.H., Estess P.,
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                     Mammalia; Eutheria;
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immunoglobulin heavy
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15-JUL-1999
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HEAVY CHAIN V REGION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EVQLQQSGAELVRAGSSVKMSCKASGYTFTSYGINWVKQRPGQGLEWIGYINPGNGYINY
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. 01, Last sequence 38, Last annotation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       try is copyright. It is produced through a collaboration -
                     Chordata;
Primates;
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QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60

Query Match Best Local

Similarity

29.8%; 65.0%;

Score 384.5; Pred. No. 5.6

.6e-23;

DB

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Length

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PIR; A02035,
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                                                                                                                                            heavy chain constant region domains.";
Biochemistry 21:5415-5424(1982).
-i- MISCELLANEOUS: THE SEQUENCE OF THE LIGHT CHAIN.
                                                                                                                                                                                                                                                                                                                        21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
IG HEAVY CHAIN V REGION MOPC 104E.
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-i- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
PIR; A01887; K2HUML.
                                                                                                                                                                                                                      MEDLINE-83075344; PubMed-6816276; Kehry M.R., Fuhrman J.S., Schilling
                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                   Mammalia;
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                                                                                                                                                                                           'Complete amino acid sequence of a mouse
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                                                                                                                    PROTEIN HAS ALSO BEEN DETERMINED.
MISCELLANEOUS: THIS PROTEIN BINDS DEXTRAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SGVPNRESGS3SGTBFTLKISRVZAZBVGVYYCMQALQTPLTFGGGTNVEIK
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                                                                                                                                                                                                                                                                                 Eutheria;
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55.
117
117 AA;
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2 AA;
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                                                                                                                                                                                                                                                                                Chordata;
Rodentia;
      12983 MW;
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                                             Glycoprotein.
BY SIMILARITY
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Pred. No. 6.
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BY SIMILARITY.
                               N-LINKED (GLCNAC. .
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                                                                                                                                                                                                                                                                                 Craniata; Ver
Sciurognathi;
    3CF8ACE4BE447E41 CRC64;
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ion update)
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thi; Muridae;
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                                                                                                                                                                                                                                                                                               Euteleostomi;
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                                                                                                                                                                                                                                                                                 Murinae;
                                                                                                                                                 THIS IGM MYELOMA
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RESULT 14

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(In) Bach M.K. (eds.);
Immediate hypersensitivity: modern concept
Marcel Dekker, New York (1978).
-1- MISCELLANEOUS: THIS EPSILON CHAIN WAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human)
Eukaryota; Metazoa; Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence upd
15-JUL-1999 (Rel. 38, Last annotation u
15-HEAVY CHAIN V-I REGION ND PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PFAM; PF00047;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. MEDLINE=83065234; PubMed=6815656;
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                                                                                                                                          GTTVTVSS
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                                                                                                                                                                                  GTTVTVSS 116
                                                                                                                                                                                                                                            APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC---
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                                                                                                                                                                                                                         APRFQGRVTMTRDASFSTAYMDLRSLRSDDSAVFYCAKSDPFWSDYYNFDYSYTLDVWGQ 135
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IH -> HI (IN REF. 2).

VG -> GV (IN REF. 2).

MISSING (IN REF. 2).
                                                                                                                                                                                                                                                                                                                                                                                                       Score 384;
Pred. No. 7
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Catarrhini; Hominidae
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R (FRAGMENTS).
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a myeloma
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Eukaryota; Metazoa; Chordat
  DOMAIN
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Appella E.;
"Amino acid sequence of the light chain variable
                                                                                                                                                                                                                                                                                                                                                                    Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota;
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IISCELLANEOUS: THIS CHAIN
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Eutheria; Rodentia;
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Eutheria; Rodentia;
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region;
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38, Last annotation update)
II REGION MOPC 511.
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COMPLEMENTARITY-DETERMINING 1
FRAMEWORK 2.
COMPLEMENTARITY-DETERMINING 2
FRAMEWORK 3.
COMPLEMENTARITY-DETERMINING 3
FRAMEWORK 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 382; DB
Pred. No. 8.4e
L5; Mismatches
                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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FT DOMAIN 40 54 FRAMEWORK 2.

FT DOMAIN 55 61 COMPLEMENTARITY-DETERMINING 2.

FT DOMAIN 94 102 COMPLEMENTARITY-DETERMINING 3.

FT DOMAIN 103 112 FRAMEWORK 4.

FT DISULFID 23 93 BY SIMILARITY.

FT DISULFID 23 93 BY SIMILARITY.

FT DISULFID 23 93 BY SIMILARITY.

FT DOMAIN 103 113 SQ SEQUENCE 113 AA; 12496 MW; 42C019D10ADA3C91 CRC64;

Query Match 29.6%; Score 382; DB 1; Length 113; Best Local Similarity 67.0%; Pred. No. 8.4e-3; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 20; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 20; Gaps 0; Gaps 0; Matches 75; Conservative 13; Mismatches 24; Indels 20; Gaps 0; Gaps 0; Gaps 20; G
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Maximum Match 100%
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Perfect score:
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2: sp_bacteria
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Copyright (c) 1993 - 2000 Comp
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ALIGNMENTS

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 kfqgkatmtadtssntaylqls	QVQLQQSGAELVRSGASVKLSCTAS(: : : : VKLQQSGGG;JVKPGGSLKLSCAAS(54.0%; Similarity 53.3%; 1; Conservative 43	•	FRAM; PF00047; ig; 2. SEQUENCE 298 AA; 31867 MW;	HSSP; P01607; TREI.	(DEC-1999) to the 36341; BAA88633.1;	method.";	ion of a novel type of	ra N., Demura T., Fuku	SEQUENCE FROM N.A.	Submitted (DEC-1999) to the EMBL/GenBank/DDBJ	STRAIN=BALB/C; TISSUE=SPLEEN; Shinohara N., Demura T., Fukuda	SEQUENCE FROM N.A.	NCB1_Tax1D=10090;	Mammalia; Eutheria; Rodentia;	Eukaryota; Metazoa; Chordata;	Mus musculus (Mouse).	FV.	0 (TrEMBLrel. 15,	(TrEMBLiel. 13,	(TrEMBLrel. 13,	PRELIMINARY;	•: • .
APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCNFYGDALDYWGQGTTVTVSSG	FNIKDYYMHWVKQ : : : SDFSRYWMSWVRQ	Score 696.5; DB 11; Pred. No. 2.9e-54; 43; Mismatches 63; I		E0F96B8A17004317 CRC64;		EMBL/GenBank/DDBJ databases	Otaticy dsing a phage	"Isolation of a novel type of vascular cell wall-specific monoclonal	da H.;		MBL/GenBank/DDBJ databases.	da H.:			Sciurognathi; Muridae;	Craniata; Vertebrata;				Last sequence update)	Created)	PRT; 298 AA.	
DYWGQGTTVTVSSG 117	RPEOGLEWIGWIDPENGDTEY 60 : : :: APGKGLEWIGEINPDSSTINY 99	Length 298; Indels 9; Gaps 3;		64;		ases.	buade arsbray subcraction	ecific monoclonal			ases.				; Murinae; Mus.	Euteleostomi;							
YWGQGTTVTVSSG 117	IGWIDPENGDTEY 60	h 298; 9; Gaps		4;		ses.	inspray subcraction	icific monoclonal			ses.				Murinae; Mus.	Euteleostomi;							

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EMBL; AF035034; AAD567
INTERPRO; IPR003006;
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01-MAY-2000 (TrEMBLrel. 14, Last sequence update)
01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
SEQUENCE FROM N.A. STRAIN-BALB/C; Malkiel S., Liao I
                                                                                        01-OCT-2000 (TrembLrel. 15, 01-OCT-2000 (TrembLrel. 15, 01-OCT-2000 (TrembLrel. 15, ANTI-MYOSIN IMMUNOGLOBULIN
                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
Wu X., Liu B., Van der Merwe P.L.,
                                                                                 Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                PFAM; PF00047;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Young D
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Primates;
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Rodentia;
  Cunningham
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. Last sequence update)
. Last annotation update)
. HEAVY CHAIN VARIABLE REGION (FRAGMENT).
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Pred. No. 1.2e
ll; Mismatches
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Sciurognathi; Muridae;
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; Murinae; Mus
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Q9Z1C4;
Q9Z1C4;
Q1-MAY-1999
Q1-MAY-1999
Q1-MAY-2000
    O90XE9,
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
IMMUNOGLOBULIN HEAVY CHAIN V-D-J REGION (FRAGMENT).
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                                                                 Q9QXE9
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STRAIN-BALB/C;
STRAIN-T.P., Giannoni M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAY-1999 (TIEMBLIEL 10, Created)
01-MAY-1999 (TIEMBLIEL 10, Last sequence update)
01-MAY-2000 (TIEMBLIEL 13, Last annotation update)
01-MAY-2000 (TIEMBLIEL 13, Last annotation update)
ANTI-PORCINE VCAM MAB 3F4 HEAVY CHAIN VARIABLE REGION
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IgG2/G4 constant regions
endothelial cells.";
                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                      PFAM;
                                                                                                                                                                                                                                                                                                                        EMBL; U78801; AAD00293.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10090;
                                                                                                                        61
                                                                                                                                           61
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                                                                                                                                                                              1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9
                                                                                        G
                                                                                                                                   APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC--NFYGDALDYWGQGTTVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEYAPKFQGKA
                                                                                                                        TQKFRGKATLTADKSSSTAYMQLSSLASEDSAVYYCARRTVGGYFDYWGQGTTLTVSS
                                                                                                                                                                   QVQVQQSGAELARPWASVKLSCKASGYNFNSYWMQWVKQRPGQGLEWIGAIYPGDGDTSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITSDTSSNTAXLQLSSLTSEDTAVXYCVRRGAVVFDYWGQGTALTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TMTADTSSNTAYLQLSSLASEDTAVYYCNFYGDAL-DYWGQGTTVTVSS 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AELVKPGASVKLSCTASGFNIEDTYMHWVKQRPEQGLEWIGRIDPATGHSKYDPKFQGKA
                                                                                                                                                                                                                                                                                                 PF00047; ig; 1.
                                                                                                                                                                                                               84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86;
                                                                                                                                                                                                                                                                                                                                     ed (NOV-1996) to the
                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity .
                                                                                                                                                                                                                                                                                                              IPR003006;
                                                                                                                                                                                                                                                                  1:
118 ·
118 AA;
                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                  118
13036 MW;
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                                                                                                                                                                                                                       33.2%;
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78.98;
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                                                                                                                                                                                                                                                                                                                                     EMBL/GenBank/DDBJ
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                                                                                                                                                                                                                       Score 428;
Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                          Hartman
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Pred. No.
                                                                                                                                                                                                                                                                     90EEC559D31EC4FC CRC64;
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                                                                                                                                                                                                             Mismatches
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No. 2.6e-32;
                                                                                                                                                                                                                                                                                                                                                       clonal antibodies with chimeric leukocyte binding to porcine
                                                                                                                                                                                                                                                                                                                                                                                          S.L.,
                                                                                                                                                                                                                        DB 11;
.9e-31;
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(Mouse)

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RESULT
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ID Q9
AC Q0
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DT 00
DT 00
DT 01
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Best Local S
Matches 78
                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 77; Conser
                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.

Clemens A., Rademaekers A., S
Submitted (DEC-1997) to the E
EMBL; AJZ25171; CAB65236.1; -
INTERPRO; IPRO03006; -.
PFAM; PF00047; ig; 1.
              Q9UL95;
01-MAY-2000
01-MAY-2000
01-JUN-2000
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01-MAY-2000
01-JUN-2000
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NON_TER
SEQUENCE
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NON_TER
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                                                             Q9UL95
                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UN-2000 (TrEMBLrel. 14, Last annotation update)
IMMUNOGLOBULIN HEAVY CHAIN V-D-J REGION (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eŭkaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse).
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                                                                                                                             APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC----NFYGDALDYWGQGTTYTVSS 116
                                                                                                                                                                          QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
                                                                                                                 NQKFKGKATLTVDKSSSTAYMQLNSLTSEDSAVYYCARDKDYY---FDYWGQGTTLTVSS
                                                                                                                                                     PF00047; ig; 1.
ER 1 1
ER 117 117
NCE 117 AA; 13000 MW;
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117
117
              (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                             PRELIMINARY
MBLrel. 13, Created)
MBLrel. 13, Last sequence update)
MBLrel. 14, Last annotation update)
IMMUNOGLOBULIN HEAVY CHAIN VARIABLE
                                                                                                                                                                                                                                                                   117
13060
                                                                                                                                                                                                                                                                                                                                                                                              Chordata;
Rođentia;
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Rodentia;
                                                                                                                                                                                                                    30.8%;
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                                                                                                                                                                                                                                                                    MW;
                                                                                                                                                                                                             15;
                                                                                                                                                                                                          Score 396.5;
Pred. No. 4.3e
L5; Mismatches
                                                                                                                                                                                                                                                                                                                                      Specht C., Koelsch E.;
EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Specht C., Koelsch EMBL/GenBank/DDBJ (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 401.5; DB 11;
Pred. No. 1.5e-28;
5; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDDE2AF84D499734 CRC64;
                                                                                                                                                                                                                                                                   D816AD0858A47E4C CRC64;
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                                                           125
                                                                                                                                                                                                          5; DB 11;
4.3e-28;
hes 21;
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   REGION
   (FRAGMENT).
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Best Local S
Matches 76
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Best Local S
Matches 74
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Eukaryota; Metazoa; Cl
Mammalia; Eutheria; Pu
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                              Q9JL82;
Q9JL82;
Q1-OCT-2000
Q1-OCT-2000
Q1-OCT-2000
                                                                                                                                              NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                              Malkiel S., Liao L., Cunningham M.W., Diamond B.; "Characterization of cross-reactive monoclonal anti-myosin/anti-n-acetyl-glucosamine antibodies from mice with autoimmune myocardit: Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     [1]
SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                     01-0CT-2000 (TremBirel. 15, Created)
01-0CT-2000 (TremBirel. 15, Last sequence update)
01-0CT-2000 (TremBirel. 15, Last annotation update)
aNTI-MTOSIN IMMÜNOGLOBULIN LIGHT CHAIN VARIABLE REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NON_TER
                                                                                                                                                                                                                                      SEQUENCE FROM N'A.
STRAIN-BALB/C;
                                                                                                                                                                                                                                                                         NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INTERPRO; IPR003006;
PFAM; PF00047; 1g; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fetus."
                      200
                                                                   140
                                                                                                                                                                                                                                                                                                                                                                                                                                                            112
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 61
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           GSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGTKLEIK
                                                      LTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRPGQSPKRLIYLVSKLDSGVPDRFT 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
                                            LSLPVSLGDQASISCRSSQSLVHTNGNTYLHWYLQKPGQSPKLLIYKVSNRFSGVPDRFS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APKFQGKATMIADTSSNTAYLQLSSLASEDTAVYYC-----NFYGDALDYWGQGTT 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EVQLVESGAEVKKPGASVKVSCKASGYTFTGYYMHWVRQAPGQGLEWMGWINPNSGGTNY
GSGSGTDFTLKISRVEAEDLGVYFCSQTTHVPYTFGGGTKLEIK
                                                                                                                                                                          AF206024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      h 30.5%;
Similarity 59.2%;
74; Conservative 1
                                                                                         Similarity 73.176; Conservative
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125
125 AA;
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                                                                                                                                               104 AA;
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                                                                                                                                               11360 MW;
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Primates;
                                                                                                   30.5%;
73.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16;
                                                                                         12;
                                                                                        Score 393; DB 11;
Pred. No. 7.6e-28;
2; Mismatches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 393.5; DB 4; Pred. No. 8.7e-28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                               5DA8BBFD5F0AA1AE CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0D3CD5C232488EAC CRC64;
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                                                                                         16;
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                                                                                                             Length
104
                     243
                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                      (FRAGMENT).
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RESULT POPULS AC 050 POPULS AC 050 POPULS OC 050 POPULS AC                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT
Q9UL92
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Best Local :
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    NON_TER
NON_TER
SEQUENCE
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NON_TER
SEQUENCE
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01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
01-JUN-2000 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
Wu X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota;
Mammalia; 1
                                                                                                                                                                                                            Wu Х.,
                                                                                                                                                                                                                                                                                                                                                                              MYOSIN-REACTIVE IMMUNOGLOBULIN HEAVY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9UL94
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                                                               РБАМ; PF00047;
                                                                              INTERPRO;
                                                                                                        EMBL; AF035020; AAD56256.1;
                                                                                                                                                   fetus.
                                                                                                                                                                                         Young D.C.;
                                                                                                                                                                                                          MEDLINE=98277139; PubMed=9614934; Wu X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                         NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                Mammalia;
                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human)
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1. 13, Last sequence update)
1. 14, Last annotation update)
1. 14, HEAVY CHAIN VARIABLE R
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13, Last sequence update)
14, Last annotation update)
10 PRILIN HEAVY CHAIN VARIABLE
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    MW;
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                                                                                                                           87:184-192(1998).
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Pred. No. 1.8e-27
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                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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    13E64F5345F4A16E CRC64;
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Q9JL75;
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Q9JL81;
01-OCT-2000
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SEQUENCE
EMBL;
                       Malkiel S., Liao L., Cunningham M.W., Diamond B.;
"Characterization of cross-reactive monoclonal anti-myosin/anti-n-acetyl-glucosamine antibodies from mice with autoimmune myocarditi submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
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"Characterization of cross-reactive monoclonal anti-myosin/anti-n-
"Characterization of cross-reactive monoclonal anti-myosin/anti-n-
acetyl-glucosamine antibodies from mice with autoimmune myocarditis.";
submitted (NOV-1999), to the EMBL/GenBank/DDBJ databases.
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
ANTI-MYOSIN IMMUNOGLOBULIN HEAVY CHAIN VARIABLE REGION
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01-OCT-2000
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Mammalia; Eutheria; Rodentia;
                                                                                                                       STRAIN-BALB/C;
                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                  NCBI_TaxID=10090;
                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                       ANTI-MYOSIN IMMUNOGLOBULIN
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114 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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12829 MW;
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Rodentia;
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                                                                                                                                                                                                                                                                                                    , Last sequence update)
, Last annotation update)
HEAVY CHAIN VARIABLE REGION
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Pred. No. 5.4e
L5; Mismatches
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No. 5
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No. 4.:
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.4e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .2e-27
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                                                   myocarditis.";
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Best Local S
Matches 72
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Best Local Similarity
Matches 73; Conser
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STRAIN-DBA/2;

Malkiel S., Liao L., Cunningham M.W., Diamond B.;

Malkiel S., Liao L., Cunningham M.W., Diamond B.;

"Characterization of cross-reactive monoclonal anti-myosin/anti-n-
acetyl-glucosamine antibodies from mice with autoimmune myocarditi-
submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.

EMBL; AF206029; AAF69327.1; -.
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NON_TER
SEQUENCE
[1]
SEQUENCE FROM N.A.
MEDLINE-98322155; PubMed-9657749;
MEDLINE-983622155; PubMed-9657749;
                                                                                                                                                                                                                                                            Q9Y298 PRELIMINARY;
Q9Y298;
01-NOV-1999 (TrEMBLrel.
01-NOV-1999 (TrEMBLrel.
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Q9JL77;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
01-OCT-2000 ITEMBLE. 15, Last annotation update)
01-OCT-2000 IMMUNOGLOBULIN HEAVY CHAIN VARIABLE REGION (FRAGMENT).
                                                                                                                                                                                                                 01-NOV-1999 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
IGG VH PROTEIN PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse).
                                                                                                               SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    71 TADTSSNTAYLQLSSLASEDTAVYYC----NFYGDALDYWGQGTTVTVSS 116
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72; Conservative
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110 AA;
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                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
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12138 MW;
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12, Last sequence update)
13, Last annotation update)
                                                                                                                                                                                                                     (FRAGMENT)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 372; DB 1
Pred. No. 6e-26;
                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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).4e-27;
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Best Local S
Matches 71
                                                                                                        Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                      Q921C6 PRELIMINARY; PRT; 117 AA.
Q921C6;
Q921C6;
Q1-MAY-1999 (TrEMBLrel. 10, Created)
Q1-MAY-1999 (TrEMBLrel. 10, Last sequence update)
Q1-MAY-2000 (TrEMBLrel. 13, Last annotation update)
ANTI-PORCINE VCAM MAB 2A2 HEAVY CHAIN VARIABLE REGION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NON_TER
SEQUENCE
                                                                                                                                                                    NON_TER
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                             Mueller J.P., Giannoni M.A., Hartman Matis L.M., Evans M.J.; Watis L.M., Evans M.J.; Humanized porcine VCAM-specific monc IgG2/G4 constant regions block human endothelial cells."; Submitted (NOV-1996) to the EMBL/Genform
                                                                                                                                                                                                      EMBL; U78799; AAD00291.1; INTERPRO; IPR003006; -
                                                                                                                                                                                                                                                                                                     STRAIN-BALB/C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Mechanism and kinetics of factor VIII inactivation: study with an IgG4 monoclonal antibody derived from a hemophilia A patient with inhibitor.":
                                                                                                                                                        SEQUENCE
                                                                                                                                                                                         PFAM; PF00047; ig; 1.
                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=10090;
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                                             QYQLQQSGPQLVRPGTSVKISCKASGYSFTSYWMHWVKQRPGQDLEWIGMIDPSDSEVKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QVQLVQSGAEVKKPGASVKVSCKVSGYTLTELPVHWVGQAPGKGLEWVGSFDPESGESIY
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NQRLKDKAILTVDKSSNTAYMQFSGPTSEDSAVYYCT-RGEVSWFAYWGQGTLVTVSA
          ch 27.5%;
l Similarity 58.5%;
69; Conservative 1
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117 AA;
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150 AA;
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                                                                                                                                                        13122 MW; 4F65B193AFB77E5B CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28.5%;
60.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 367.5; DB '
                                                                                            Score 354.5; DB 1
Pred. No. 2.3e-24;
5; Mismatches 31
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                                                                                                                                                                                                                                                                    monoclonal antibodies with chimeric
                                                                                                                                                                                                                                                       leukocyte
                                                                                                                                                                                                                                                                                         S.L., Elliott E.A.,
                                                                                                                     DB 11; Length
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Search completed: February 12, 2001, 15:24:31 Job time: 101 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum DB seq
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1: /SIDS1/gcgda
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Gapop 10.0 , Gapext 0.5
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W28491
W28492
Y06273
W60770
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294.127 Million cell updates/sec
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                                               Single chain antib
Human p53 protein
Human p53 protein
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Single chain antib
Fusion protein pIC
Fusion protein (80 :
MFE-23 antibody.
Single chain anti-
             Biosynthetic antib
                                                                                                                                                                                                                                             Description
                                   H22-anti-CEA antib
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RESULT W60769

W60769 standard; Protein; 243

AA.

W60769;

08-SEP-1998 (first entry)

Single chain antibody (ScFv) 421 that binds to mutant p53 proteins.

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13
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W89101	W95600	W95462	Y28470	R21262	Y87477	W37132	W90218	R72599	Y17965	W35132	W35131	W90217	W27560	Y51142	Y59265	Y17960	R95213	R32090	Y17964	R32091	W63830	W82742	Y06132	W82743	P80151	Y21884	537	796	1795	991		3
Anti-digo	Anti-digoxin	Anti-Tac(Vh-(Lk)-V	pscFvNQ11	Anti-Tac(Anti-Tac(Bispecific	Anti-dans	Mouse sci	R. pipiens	R. pipier	Bispecific	Consensus	Murine de	Antibody		an	H and L c	Mouse scr	H and L c	Transformant	Plasmid p	Monoclona	Fusion pr	Multifunctional	Amino acid	Ė	Ø	Mouse sci	Human MCP	ש	Bispecifi
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                                              WPT; 1998-272149/24.
N-PSDB; V36236.
                                                                                                                                                                                                                                                                                                                                           Single chain antibody; ScFv 421; mouse; p53 protein; oligomerisation; regulatory domain; p53 mutant; H273; W248; G281; p53-dependent trans-activating activity; restoration; tumour-suppressing activity; tumour_cell; treatment; hyper-proliferation; cancer; re-stenosis; ss.
Restoring p53-dependent trans-activating activity to cell containing mutant p53 - by delivering single-chain antibody specific for the
                                                                                                                                                                           29-OCT-1996;
                                                                                                                                                                                                         27-OCT-1997;
                                                                                                                                                                                                                                           07-MAY-1998
                                                                                                                                                                                                                                                                                                             Mus sp.
                                                                                                      Debussche L,
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                                                                                                                                          (RHON ) RHONE-POULENC RORER SA
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W37738

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a single chain antibody (ScFv) designated 421. The antibody binds to an epitope present in the C-terminal region of the p53 protein that includes oligomerisation and regulatory domains, specifically between positions 320 and 393. ScFv 421 is directed against p53 mutants, particularly H273, W248 and G281 mutants. When the ScFv is introduced into cells containing a mutant p53 protein, p53-dependent trans-activating activity is restored. ScFv 421 is specific for p53-mutants that have lost tumour-suppressing activity and are present in tumour cells. It is particularly used to treat hyper-proliferation associated with these mutants (e.g. cancer and re-stenosis) but may also be used in vitro for studying mechanisms of activity of p53 or its mutant and to purify or detect p53.
                            19-JUL-1995;
                                                       17-JUL-1996;
                                                                                                                WO9704092-A1
                                                                                                                                                        Chimeric - Homo sapiens.
Chimeric - Herpes simplex virus
                                                                                                                                                                                                    tumour suppression;
                                                                                                                                                                                                                Leucine zipper domain; LZD; oligomerisation domain; mutant; mute substitution; replacement; transactivation; viral protein VP16; anti-oncogene; hyperproliferation; cancer; restenosis; ScFv;
                                                                                                                                                                                                                                                                       Human p53 protein variant S-325 encoded by pEC176.
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(RHON *) RHONE
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POULENC RORER SA
                            95FR-0008729
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                                                                                                                                                                                                    apoptosis; single chain
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Pred. No. 2e-83;
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Best Local S
Matches 240
                            *Leucine zipper domain; LZD; oligomerisation domain; mutant; muta
substitution; replacement; transactivation; viral protein VP16;
anti-oncogene; hyperproliferation; cancer; restenosis; SCFV;
tumour suppression; apoptosis; single chain antibody variable do
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claimed variants of protein p53 have at least part of the coligomerisation domain deleted and replaced by a leucine zipper domain. The mutants preferably also have at least part of the p53 transactivation domain (amino acids 1-74) deleted and replaced by the transactivating domain (TD) from herpes simplex virus viral protein vP16 (amino acids 411-490) or by a protein domain able to bind selectively to a transactivator, especially a single-chain antibody variable domain (ScFv). The present sequence is that of a specifically claimed p53 variant designated S-325 and comprising a ScFv domain, amino acids 75-325 of human wild-type p53 and a leucine zipper domain at the C-terminal. The p53 variants are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              more active and more stable tumour suppressors and apoptosis-inducing agents than wild-type p53 and are active where the wild-type protein is not, i.e. they are not inactivated by dominant negative or oncogen mutants, nor by other cellular proteins (because the leucine zipper domain prevents formation of inactive mixed oligomers).
                                                                                                           Human
Chimeric -
                                                                                                                                                                             W28492;
                                                                                                                                                                                                             W28492 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                             25-NOV-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           esp. cancer and restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New p53 variants e.g. with oligomerisation domain replaced by leucine zipper - useful for treating hyper-proliferative disorders,
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                                                                                                             p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T86221.
                                                                                                           protein variant S-325H.
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conseiller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  535 AA;
                                                                                                                                           (first entry)
                                                                                                                                                                                                             Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99.1%;
                                                                                                                                                                                                              535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 1277; DB 18
Pred. No. 3.1e-82;
2; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      535;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or oncogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                mutein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60
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                                                                                                                                                                                                                                                                                                                                                                                          240
                                                                                                                                                                                                                                                                                                                                                                                                                          182
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Claimed variants of protein p53 have at least part of the oligomerisation domain deleted and replaced by a leucine zipper domain. The mutants preferably also have at least part of the p53 transactivation domain (amino acids 1-74) deleted and replaced by the transactivating domain (TD) from herpes simplex virus viral protein vP16 (amino acids 411-490) or by a protein domain able to bind selectively to a transactivator, especially a single-chain activated and replaced by the selectively to a transactivator, especially a single-chain activated as specifically claimed p53 variant designated s-325H and comprising a scFv domain, amino acids 75-325 of human wild-type p53 (but with Arg182 replaced by His) and a leucine zipper domain at the C-terminal. The p53 variants are more active and more stable tumour suppressors and apoptosis-inducing agents than wild-type p53 and are active where the wild-type protein is not, i.e. they are not inactivated by dominant negative or oncogenic mutants, nor by other cellular proteins (because the leucine zipper domain prevents formation of inactive mixed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 36; Page -; 133pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New p53 variants e.g. with oligomerisation domain replaced by leucine zipper - useful for treating hyper-proliferative disorders, esp. cancer and restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RHON ) RHONE POULENC RORER SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                       241
                                                                        183
                                                                                                                      181
                                                                                                                                                                       123
                                                                                                                                                                                                                     121
                                                                                                                                                                                                                                                                      63
                                                                                                                                                                                                                                                                                                                     61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 98.
                                                                                                                                                                                                                                                                                                                                                              e: this sequence does not appear in the specification and has
produced by modifying the given sequence of variant V-325).
                                                                                            KRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGTKL 240
                       EIK 243
                                                                     krliylvskldsgvpdrftgsgsgtdftlkinrveaedlgvyycwqgthspltfgagtkl
<u>..</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conseiller E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                535
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95FR-0008729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96WO-FR01111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Arg residue at position 182 of wild-type
    p53 has been mutated to His"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99.1%;
98.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 1277; D
Pred. No. 3.1e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 18;
3.1e-82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 535;
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         The present sequence represents a fusion protein comprising murine canti-Fc alpha raceptor antibody A77 sFv and the transmembrane domain of platelet derived growth factor receptor (PDFR-R TM). It is encoded by expression vector pJG718 (see X5935). Murine tumour cells transformed to express A77-TM were able to bind a soluble form of the Fc alpha receptor, and the bound receptor was able to engage CI GA molecules. This is an example of cells of the invention that the been transformed to express on their surface a component which binds to an Fc receptor of an effector cell. The transformed cell is targeted to an effector cell via the Fc binding component, and can be used asia vehicle to increase an effector cell-mediated antigen associated with the cell. The transformed cells are used to treat cancer and infectious diseases or used as vaccines. The method allows for killing of target cells without targeting any particular antigen on the cell. This is advantageous since many
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Deo YM, Goldstein J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single chain antibody; scFv; A77; IgA receptor; Fc receptor; Fc alpha receptor; platelet derived growth factor receptor; antibody engineering; cell surface expression; therapy; cancer; tumour; vaccine; human.
                                                                                                                                                                                                                                                                                                                          Example 6; Fig 10A-E; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                Cells expressing anti-Fc receptor binding components
                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-371099/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-AUG-1999 (first entry)
tumour cells and other target cells do not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        receptor scFv A77-PDGR-R TM fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0067232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US25556
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176..287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note= "A77 VL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "HA epitope"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Graziano R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "PDGFR transmembrane domain"
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The present s
D3M. The anti
of the p53 pro
specifically l
                                                    Claim
                                                                    Restoring p53-dependent trans-activating activity to cell mutant p53 - by delivering single-chain antibody specific mutant, particularly for treatment of tumours
                                                                                                                                                                                                                             07-MAY-1998
                                                                                                                                                                                                                                                                                     p53-dependent trans-activating activity; restoration; tumour-suppressing activity; tumour cell; treatment; hyper-proliferation; cancer; re-stenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                            W60770
                                                                                                                                                                                                                                                                                                                                                                                                              W60770
                                                                                                                                                                                     29-OCT-1996;
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                                                                                                                                                               (RHON ) RHONE-POULENC
                                                                                                                                                                                                                                                                                                                     regulatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match
present sequence represents a single chain an I. The antibody binds to an epitope present in the p53 protein that includes oligomerisation orifically between positions 320 and 393. ScFv
                                                                                                                                                                                                                                                                                                                                                                                                                                  <sub>5</sub>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VSSGGGSGGGSGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            eiqlqqtgpelvkpgasvkisckasgysftdylifwvkqshgkslewtgnnnpyygstsy
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fgggtkleik
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vsagggggggggggggggdvvmtqtpltlsitigqpasisckssqslldsdgktylnwll
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC----NFYG---DALDYWGQGTTVT 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       188;
                                                    'n
                                                                                                                                                                                                                                                                                                                    chain antibody; ScFv D3M; mouse; p53 protein; oligomerisation;
tory domain; p53 mutant; H273; W248; G281;
                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                    chain
                                                    Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                              Protein;
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                                                                                                                                                                 RORER
                                                                                                                                                                                                                                                                                                                                                   (ScFv)
                                                    French
                                                                                                                                                                                                                                                                                                                                                                                                               249
                                                                                                                                                                 SA
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Pred. No. 5.2e
23; Mismatches
                                                                                                                                                                                                                                                                                                                                                  D3M that binds to mutant p53 proteins
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 n antibody (ScFv) do
in the C-terminal :
ion and regulatory of
CFv D3M is directed
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 directed
                                                                                containing for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           353;
                              designated
           domains,
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                    region
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                                        New-yene construct expressing conjugate of targeting a prodrug-converting enzyme - useful for, e.g. targeted cytotoxic drug in vivo, especially for treatment of converting in vivo, especially for treatment of converting the statement of converting the st
                                                                                                                                                                                                         Blakey DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conjugate; cell targeting; cytotoxic drug; plasmid; fusion protein; prodrug-converting enzyme; cell surface antigen; treatment; cancer; inflammation; rheumatoid arthritis; antibody; prodrug therapy syste
                                                                                                                                              N-PSDB;
                                                                                                                                                                                                                                                                                                                       10-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fusion
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Pred. No. 1.9e
23; Mismatches
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Best Local S
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                                                                                                                                                                                                                                                                Conjugate; cell targeting; cytotoxic drug; plasmid; fusion protein; prodrug-converting enzyme; cell surface antigen; treatment; cancer; inflammation; rheumatoid arthritis; antibody; prodrug therapy syste
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence is a used in a method for obtaining a novel gene construct (A) which expresses, in cells of a mammal, a conjugate (B) of a cell-targeting group (I) and a heterologous prodrug-converting enzyme (II), and (B) is directed to leave the cell for selective localisation
                           N-PSDB;
                             WPI; 1999-059700/05
N-PSDB; V72075.
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gtkleik 263
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construct expressing conjugate of targeting agent
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                                                                     Emery SC;
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Pred. No. 6.8e-57;
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Best Local Similarity

Matches 174; Conserv
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                                                                                                                                               Кеу
                                                                                                                                                                                                                                                                                                      R75719 standard; Protein; 270
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                                                                                                                                                                        Mus musculus
                                                                                                                                                                                                           Antibody; MFE-23; carcinoembryonic antigen; CEA; colorectal tumour;
                                                                                                                                                                                                                                    MFE-23 antibody.
                                                                                                                                                                                                                                                           26-JAN-1996 (first
                                                                                                                                                                                                                                                                                     R75719;
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                                                                                                                                                                                              therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGGSGGGGSGGGSDVLMTQTPLTLSVTTIGQPASISCKSSQSLLDSDGKTYLNWLLQRP 176
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                                                                                                                                                                                                 diabody.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 87-89;
62..75
/note-
                                                         /note=
27..51
                                                                                  /note=
162..2
                                                                                                           /note=
27..146
                      52..61
/note=
                                                                                                                                  Location/Qualifiers 27..267
                                              /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      70.7%; Score 911; 70.4%; Pred. No. 1.
                                                                                     . 267
                                                                                                                       "variable region"
                       "CDR H1
                                               "FR H1"
                                                                                              "variable heavy chain"
"FR H2"
                                                                       "variable light chain"
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                       Qy
                                                                                                            cancer imaging and therapy. The MFE-23 antibody sequence was obtained using phage technology. In this process, mice were immunised with CEA.

The antibody variable region genes obtained from these mice were then amplified from cDNA and cloned as a single chain Fv (scFv) into bacteriophage vectors, producing a library. The phages that bound to bacteriophage vectors, producing a library. The phages that bound to be intinylated CEA were selected and amplified, and this sequence (and the encoding cDNA) were selected. This sequence was found to have good specificity and affinity for CEA, meaning that it can be used in targeted anti-tumour therapies. A humanised antibody with the complementarity determining regions of MFE-23 may be produced by CDR grafting. The antibody may be used for the treatment by surgery or therapy of a colorectal tumour. WFE-23 may also be used to make diabodies (bivalent or bispecific antibody fragments which bind to two different antigens), and may be linked to an antibody and may be linked to an antipour agent or adetertable label.
                                                 Matches 173;
                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                   This sequence represents the MFE-23 antibody, which is an antibody against carcinoembryonic antigen (CEA). CEA is a marker antigen f
                                                                                                                                                                                                                                                                                                                                                                               Antibody for carcinoembryonic antigen - of colorectal cancer
                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                       Claim 3; Page 48-49;
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                         \vdash
1995-215234/28
                                                                                                                                  agent or a detectable label.
                                                                                                             270 AA;
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chester
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/note=
86..12
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217..2
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                                                            70.3%;
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                                                  26;
                                               Score 906; DB 16;
Pred. No. 1.5e-56;
6; Mismatches 35;
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                                                                       Length 270;
                                                 Indels 16;
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RESULT W11508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Humanised antibody; anti-Fc receptor; H22; bifunctional; bispecific; fusion protein; chimera; carcinoembryonic antigen; CEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single chain anti-Fc gamma RI antibody fused to anti-CEA antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W11508;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    W11508 standard; Protein; 553
                      WO9640789-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            174 QRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLT 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          114 VSSGGGGSGGGGGGGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLL 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              144 vssggggsggggggggsenvltqspaimsaspgekvtitcsasssv-----symhwfq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                      521..533
                                                                                                                                                                       412..520
                                                                                                                                                                                                                                                                       267..276
                                                                                                                                                                                                                                                                                                                                              140:.154
                                                                                                                                /label= MFE-23_VL
/note= "Anti-CEA antibody MFE-23
domain"
                                                                                                                                                                                               397..411
                                                                                               /label= linker
534..544
                                                                                                                                                                                                        /label- MFE-23_VH
/note- "Anti-CEA antibody MFE-23 heavy chain variable domain"
                                                                                                                                                                                                                                             277..396
                                             /note= "6 histidine residues"
                                                                                                                                                                                                                                                                                            /note= "Anti-Fc
                                                                                                                                                                                                                                                                                                                                /label= linker
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                                                                                                                                                                                  /label= linker
                                                                                                                                                                                                                                                           /label= linker
                                                                                                                                                                                                                                                                                                          /label- H22_VL
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                                                                      label= c-myc_tag
48..553
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                                                                                                                                                                                                                                                                                                                                                       H22_VH
"Anti-Fc gamma RI antibody H22 heavy chain
variable domain"
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                                                                                                                                                                                                                                                                                  e domain"
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07-JUN-1996;

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Best Local Similarity
                                                                                                  Multispecific single chain antibody; antibody H22; tumour cell; therapy; antibody-dependent cellular cytotoxicity; ADCC; HER 2/neu; infection; epidermal growth factor receptor; breast cancer; ovarian cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A mammalian expression construct encoding a single chain antibody having binding specificity for Fc gamma RI, derived from the humanised anti-Fc gamma RI monoclonal antibody H22 was prepared (see T58129). A bispecific single chain polypeptide was produced by fusing the H22 ScFv to an anti-carcinoembryonic antigen (CEA) antibody. The H22 anti-CEA fusion protein was shown to bind both
07-JUN-1996;
                         17-NOV-1998
                                                 US5837243-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-1995;
                                                                                                                                                      H22-anti-CEA antibody
                                                                                                                                                                                                                                 W73223 standard; Protein; 553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Recombinant, multi-specific anti-Fc receptor antibody molecules also comprise an anti-target portion, used for the treatment of cancer, autoimmune disease and pathogenic infection
                                                                                                                                                                               25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; Fig 40; 115pp; English
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                                                                                                                                                                                                                                                                                                                                              PKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEYA
                                                                                                                                                                                                                                                                                             gagtklelk
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                                                                                                                                                                                                                                                                                                                                                                                                              SSGGGGSGGGGSGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQ 174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         553 AA;
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                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                517
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96US-0661052.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     69.8%; Score 900; DB 18; Length 68.7%; Pred. No. 8e-56;
                                                                                                                                                     construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Graziano
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody-dependent cellular cytotoxicity (ADCC) against a tumour cell which is characterised by overexpression of HER 2/neu or epidermal growth factor receptor (EGFR), comprises contacting the tumour cell with a multispecific protein molecule (preferably a single chain antibody) comprising: (a) an anti-Fc receptor antibody or an antigen binding fragment; (b) a portion which binds to HER 2/neu; and (c) a portion which binds to DEFR. The method can be used for treating cancers especially breast cancer or ovarian cancer. The multispecific antibody can also be administered prophylactically to vaccinate a subject against infection
                           Biosynthetic antibody binding multifunctional protein.
                                                                      Biosynthetic antibody binding
                                                                                                   01-JAN-1980
                                                                                                                          P80154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   anti-CEA antibody and a multispecific single chain antibody designated H22. The antibody can be used in the method of the invention for induc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Specific killing of tumour cells - using a multi-specific molecule comprising an anti-Fc receptor antibody and a portion which binds to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1996; 1
07-JUN-1995;
W08809344-A
                                                                                                                                                          P80154 standard; protein; 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    by a target cell.
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N-PSDB; V08176.
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                                                                                                                                                                                                                       gagtklelk 517
                                                                                                                                                                                                                                                               GAGTKLEIK 243
                                                                                                                                                                                                                                                                                            kpgtspklwiystsnlasgvparfsgsgsgtsysltisrmeaedaatyycqqrssypltf 508
                                                                                                                                                                                                                                                                                                                                                                                                                            PKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                171;
                                                                                                                                                                                                                                                                                                                         RPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTF 234
                                                                                                                                                                                                                                                                                                                                                   ssggggsggggggggenvltqspaimsaspgekvtitcsasssv----symhwfqq
                                                                                                                                                                                                                                                                                                                                                                                 SSGGGGSGGGGSGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQ 174
                                                                                                                                                                                                                                                                                                                                                                                                             pkfqgkatfttdtssntaylqlssltsedtavyycnegtptgpyy---fdywgqgttvtv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       553 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                   (first entry)
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95US-0484172.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             69.8%;
                                                                                                                                                                                                                                                                                                           Graziano R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 900; DB 2
Pred. No. 8e-56;
                                                                      site.
                                         site;
                                                                                                                                                             A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                         framework region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Somasundaram
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 553;
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                                         assay;
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RESULT W37738 ID W3 XX * AC W3 XX OT 07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The biosynthetic antibody binding site forms part of a recombinant CC multifunctional protein which also comprises an effector protein, an AA cacid sequence capable of sequestering an ion, or a sequence capable of binding to a solid support. The BABS comprises the framework region from murine anti-digoxin monoclonal antibody 26-10-heavy chain variable region and the complementarity determining region from G-loop-4 heavy chain variable region and has lysozyme specificity. The effector protein is an enzyme, toxin, receptor, binding site, growth factor, cytokine or enzyme, toxin, receptor, binding site, growth factor, cytokine cor antimetabolite. The sequence capable of sequestering an ion is calmodulin or metallothionein. The sequence capable of binding to solid support is streptavidin or a protein A fragment. The protein may be used for, eg specific binding assays, affinity purificn., biocatalysts, drug targetting, imaging and immunological treatment of oncogenic etc. diseases. The protein offers fewer cleavage sites to circulating protein protein and are cleaved stability. They reach target organs regions regions and are cleared quickly from the body. They also have reduced
 07-JUL:1998
                                  W37738;
                                                                 W37738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; ; 15pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Recombinant multifunctional protein - having antibody binding sequence for biological activity, ion sequestering or binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-MAY-1988;
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                                                                                                                                                   240
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                                                                                                                                                                                                                                                                                                                                                     63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1988-353928/49
DB; N80180.
                                                                                                                                                 gtkleik 246
                                                                                                                                                                                  GTKLEIK 243
                                                                                                                                                                                                                   gqspklliykvsnrfsgvpdrfsgsgsgtdftlkisrveaedlgiyycfqgshvpwtfgg 239
                                                                                                                                                                                                                                     GQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGA 236
                                                                                                                                                                                                                                                                                  ggggsggggggggggdvvmtqtplslpvslgdqasiscrssqsivhsngntylewylqka 179
                                                                                                                                                                                                                                                                                                   GGGGSGGGGGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRP 176
                                                                                                                                                                                                                                                                                                                                                                                    APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC----NFYGDALDYWGQGTTVTVSS 116
                                                                                                                                                                                                                                                                                                                                                                                                                    evqlqqsgpelvkpgasvrmsckssgytftnyyihwlkqshgkslewigwiypgngntky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           support.
                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       249 AA;
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 (first
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                                                                   Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    69.5%; Score 895.5; 67.6%; Pred. No. 7.
                                                                   437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .3e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       249;
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to a
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Best Local
                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                               This is the amino acid sequence of a fusion protein comrising of Mgr6 single chain antibody (with the heavy and light chain joine a linker), fused to the clavin protein. Clavin is an inhibitor protein, and functions by inactivating the ribosomes. Clavin or conjugates are useful as anticancer and/or antiviral agents. The recombinant ribosomal inhibitor protein (RIP), Mgr6-clavin (a co of clavin) is used as an immunoconjugate, and the complex and cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Pages 14-15; 26pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA encoding Aspergillus clavatus-derived ribosomal protein, clavin - useful as an immuno-conjugate and
                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                   alone are capable of inhibiting ribosomal activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Colnaghi M, De Santis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-DEC-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-077109/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ITUY-) ITAL MIN UNIV RICERCA SCI & TECNOLOGICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleotide sequence encoding the Mgr6-clavin immunotoxin
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276 klelk 280
                         239 KLEIK 243
                                                    216 ppkllikyasnlesgvparfsgsgsgtdftlnihpveeedtatyycqhsweiprtfgggt
                                                                                                              157
                                                                                                                                          119
                                                                                                                                                                     97
                                                                                                                                                                                               61
                                                                                                                                                                                                                             37
                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                         ب
                                                                                                           ggsggggggggyieltqspaslavslgqratiscrasqs-vstsrysymhwyqqkpgq
                                                                                                                           GGSGGGSGGGGSDVLMTQTPLTLSYTIGQPASISCKSSQSLLDSDGKTYLNWLLQRPGQ 178
                                                                                                                                                                   dpkfqgkatitadtssntaylqlssltsedtavyycargeydypfpywgqgtsvtvssgg
                                                                                                                                                                                               APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC--NFYGDALDYWGQGTTVTVSSGG 118
                                                                                                                                                                                                                            QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
                                                                                 SPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthesis;
                                                                                                                                                                                                                                                                                                                                                         437 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Aspergillus clavatus
Mus musculus
                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ribosomal inhibitor protein; RIP; clavin; inhibition; thesis; conjugate; Mgr6-clavin; anticancer; antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Mgr6 from M. musculus"
287..437
/note= "~~
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                68.8%; Score 886.5; DB 170.2%; Pred. No. 5.6e-55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Clavin from A. clavatus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mele A,
                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                               DB 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 anticancer; antiviral agent
                                                                                                                                                                                                                                                                                      44;
                                                                                                                                                                                                                                                                                                               Length 437;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibitor for treatment of
                                                                                                                                                                                                                                                                                                                                                                                                  (a conjugate ind clavin
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bined by
                                                                                                                                                                                                                                                                                                                                                                                                                                           or its
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                                                                                                                                                                                                                                               an antibody which specifically interacts with the extracellular domain of Ct the human zeta-chain. The antibody whose CDR of a variable region is encoded by (1), is obtained by immunising a rat with jurkat cells and CC encoded by (1), is obtained by immunising a rat with jurkat cells and CC extractions with a conjugate comprising a carrier molecule and a copyride having 11 N-terminal amino acids of a rat zeta-chain. The CC enti-zeta-chain antibody is useful for the treatment and prevention of CC autoimmune diseases, immune deficiencies, T-cell malignancies, CC infectious diseases and the suppression of immune response preferably in CC order to avoid graft rejection after organ transplantation, malignancies, CC or viral infections. The antibody, and fragments of it, can be useful for the enhancement or suppression of NK-cell dependent immunity or for the treatment of NK-cell derived malignancies. It can also be useful for the CC treatment of NK-cell derived malignancies. It can also be useful for the CC treatment of the chain or eta-chain expression on NK-cells, CC T-lymphocytes or their precursors. The present sequence represents a bispecific anti-zeta-chain/anti-zecahain expression on NK-cells.
                                                                                                                                                  Query Match
Best Local Sim
Matches 179;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           autoimmune infectious
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                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a nucleic acid molecule (I) encoding at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bispecific anti-zeta-chain/anti-EpCAM antibody protein sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 9; Page 74-76; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotide, polypeptide, antibody useful for treating autoimmune disease, immune deficiencies, T-cell malignancies and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-160926/14.
N-PSDB; Z88358.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rattus norvegicus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complementary determining region; CDR; autoimmune disease; cytostatic; immune deficiency; T-cell malignancy; infectious disease; antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CONN-) CONNEX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-zeta-chain antibody 2-B-5; rat; human zeta chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunosuppressive;
   202
                                                                           142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13
                                     61 APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC--------
                                                                                                                                                                        Local Similarity
                                                                                                            1 QVQLQQSGAELVRSGASVKLSCTASGENIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
nqkfngkatltadkssstaymqlssltsedsavyfcardwhyyssyirpfaywgqgtlvt
                                                                     qvqlqqsgaelvkpgssvkisckasgytftsydmhwikqqpgnglewigwiypgngntky 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                one complementary
                                                                                                                                                                                                                                                 532
                                                                                                                                                                                                                                                                                     invention.
                                                                                                                                                    Conservative
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               67.48;
47.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                determining region (CDR) of a variable region of
                                                                                                                                                  27;
                                                                                                                                                  Score 869; DB 21;
Pred. No. 1.1e-53;
7; Mismatches 37;
                                                                                           Length 532;
                                                                                                                                                  Indels 136;
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          The present invention describes fusion proteins comprising a chemokine and a tumour antigen or HIV antigen. Specifically claimed fusion proteins comprise: (1) human monocyte chemotactic protein 3 (MCP-3) and human Muc-1; (2) human interferon induced protein 10 (IP-10) and human Muc-1; (3) human macrophage-derived chemokine (MDC) and human Muc-1; (4) human SDF-1 and human MCC and HIV gpl20; (6) human MCP-3 and HIV gpl20; (7) human MCC and HIV gpl20; and (8) human SDF-1 and HIV gpl20. The fusion proteins, and nucleotide sequences encoding them, can be used for producing an immune response, e.g. an effector T cell immune response. They can also be used for treating cancer or treating or preventing HIV infection. The fusion proteins and/or nucleotide sequences can be used in in vitro diagnostic assays, as well as in screening assays for identifying unknown tumour antigen epitopes and fine mapping of tumour antigen epitopes. The present sequence represents a fusion protein
                                                                                                                                                                                                                                                      Disclosure; Page 114-115; 142pp; English.
                                                                                                                                                                                                                                                                                                       New fusion polypeptides comprising a chemokine and
                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-551418/46.
                                                                                                                                                                                                                                                                                                                                                                                          Kwak LW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chemokine; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human IP-10 and murine scrv20A fusion protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Y29910 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                        (USSH ) US DEPT HEALTH & HUMAN SERVICES
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Matches 160
The present invention describes fusion proteins comprising a chemokine and a tumour antigen or HIV antigen. Specifically claimed fusion proteins comprise: (1) human monocyte chemotactic protein-3 (MCP-3) and human Muc-1; (2) human interferon-induced protein 10 (IP-10) and human Muc-1; (3) human macrophage-derived chemokine (MDC) and human Muc-1; (4) human SDF-1 and human Muc-1; (5) human IP-10 and HIV gp120; (6) human MCP-3 and HIV gp120; (7) human MDC and HIV gp120; and (8) human SDF-1 and HIV
                                                                                                                                  New fusion polypeptides comprising a chemokine HIV antigen, used for treating cancers or treat
                                                                                                                                                                                               Kwak LW,
                                                                                                  Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
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60; Conservative
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                                                                                                 Page 116-117; 142pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                            and murine scFv20A fusion protein
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Compugen Ltd

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ALIGNMENTS

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SOFTWARE: Patentin Ver. 2
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LENGTH: 243
TYPE: PRT
ORGANISM: Mus musculus
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EARLIER APPLICATION NUMBER: FR96/13176
EARLIER FILING DATE: 1996-10-29
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CURRENT FILING DATE: 1999-04-26
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IITLE OF INVENTION: ANTI-P53 SINGLE-CHAIN ANTIBODY FRAGMENTS AND THEIR USES
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       Conservative
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                                                                       100.0%;
                                                                Score 1289; DB 16;
Pred. No. 1.3e-102;
       0,
       Indels
                                                                                                                                Length 243;
0;
Gaps
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1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60

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FILING DATE: 17-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: SAVITZKY ESG., MARTIN F.
REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: ST95044
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                     TELEPHONE: (610) 454-3808
TELEFAX: (610) 454-3808
INFORMATION FOR SEQ ID NO: (SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: CONSEILLER, Emmanuer
APPLICANT: BRACCO, Laurent
TITLE OF INVENTION: P53 PROTEIN VARIANTS AND THERAPEUTICAL
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: FR 95/08729
FILING DATE: 19-JUL-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy
                                                                                                                                                                     TYPE: amino acids
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                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                                               Conservative
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98.8%;
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                                             2;
                                                             Score 1277;
Pred. No. 3
                                             Mismatches
                                           DB 13;
3.6e-101;
nes 1;
                                                                            Length
                                               Indels
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Best Local Similarity
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LENGTH: 257
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CURRENT FILING DATE: 1999-10-18
EARLIER APPLICATION NUMBER: 98 11 9630.6 EP
EARLIER FILING DATE: 1998-10-16
EARLIER APPLICATION NUMBER: 66/BOM/1998 INDIA
EARLIER FILING DATE: 1998-10-16
NUMBER OF SEQ ID NOS: 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Molecular Pathogenicide Mediated Plant Disease
TITLE OF INVENTION: Resistance
FILE REFERENCE: 0.47-0189P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: NAHRING, Jorg
APPLICANT: SACK, Markus
APPLICANT: MONECKE, Michael
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: LIAO, Yu-Cai
                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 257
TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICANT: EMANS, Neil
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179
              174 QRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLT
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63 NQKFKGKATLTADKSSSTAYMQLSSLTSEDSAVYYC----ARPDVWGAGTLLTVSAGAGP 118
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QRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKISRVEAEDLGVYYCWQGTHFPHT
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77.2<del>8</del>;
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Pred. No. 2.8
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nes 33;
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RESULT 4 US-09-297-181-4

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; TYPE: PRT
; ORGANISM: Mus musculus
US-09-297-181-4
NUMBER OF SEQ ID NOS: 3
SOFTWARE: PatentIn Ver.
SEQ ID NO 28
LENGTH: 238
TYPE: PRT
ORGANIZM: Mouse
US-09-559-019-28
                                                                                                                                                                                                                                                                                                   US-09-559-019-28
                                                                                                                                                                                                                                                                                                                          RESULT
                                                                                                                                                                                                                      Sequence 28, Application US/09559019
GENERAL INFORMATION:
APPLICANT: 2hu, Zhenping
APPLICANT: Witte, Larry
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SEO ID NO 4
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Best Local
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                                                                                                                                    TITLE OF INVENTION: Antibodies Specific to FILE REFERENCE: Sequence Listings 1-34 for CURRENT APPLICATION NUMBER: US/09/559,019
CURRENT FILING DATE: 2000-04-25
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EARLIER FILING DATE: 1997-10-27
EARLIER APPLICATION NUMBER: FR96/13176
EARLIER FILING DATE: 1996-10-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/297,181
CURRENT FILING DATE: 1999-04-26
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                                                                                                                                                                                                                                                                                                                                                                                     240
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                                                                                                                                                                                                                                                                                                                                                                                                                           FGAGTKLEIK 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SSGGGGSGGGGSGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDS-DGKTYLNWLL 173
                                                                                                                                                                                                                                                                                                                                                                                     FGGGTKLEIK 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QRPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLT 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QKPGQSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SSGGGGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQ 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60
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                                                                                                                                                                                KDR and Uses Thereof 381-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     239
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Query Match

72.28;

Score 930.5;

DB 19;

Length 238;

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PCT-US00-19843-11 , Sequence 11, Application PC/TUS0019843 , GENERAL INFORMATION:
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; ORGANISM: Mouse
US-09-559-019-27
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                                                                             RESULT
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APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
TITLE OF INVENTION: Antibodies Specific to KDR and Uses Thereof
FILE REFERENCE: Sequence Listings 1-34 for 381-71
CURRENT APPLICATION NUMBER: US/09/559,019
CURRENT FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 175; Conserv
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APPLICANT: Herr, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 240
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                                                                                                                                   235 LEIK 238
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                                                                                                                                                                     LEIK 243
                                                                                                                                                                                                            PKLWIYSTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPFTFGSGTK 234
                                                                                                                                                                                                                                                PKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGTK 239
                                                                                                                                                                                                                                                                                      GSGGGGSGGGSDIELTQSPAIMSASPGEKVTITCSASSSV-----SYMHWFQQKPGTS 174
                                                                                                                                                                                                                                                                                                                                                                  APKFQGKATMTADSSSNTAYLQLSSLTSEDTAVYYCNAYYGDYEGYWGQGTTVTVSSGGG
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71.7%;
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Pred. No. 1.5
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CURRENT APPLICATION NUMBER: PCT/US00/19843
CURRENT FILING DATE: 2000-07-21
PRIOR APPLICATION NUMBER: 60/145,512
PRIOR FILING DATE: 1999-07-23
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 11
LENGTH: 251
TYPE: PRT
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                        ; ORGANISM: Mus sp
US-09-523-095A-30
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US-09-523-095A-30
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                                                                                                                                        SOFTWARE: PA
SEQ ID NO 30
LENGTH: 271
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 30, Application US/09523095A GENERAL INFORMATION:
 Best Local Similarity Matches 174; Conserv
                                   Query Match
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Best Local Similarity
Matches 172; Conserv
                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/523,095A CURRENT FILING DATE: 2000-03-10 PRIOR APPLICATION NUMBER: JP 11-63557 PRIOR FILING DATE: 1999-03-10
                                                                                                                                                                               NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver.
                                                                                                                                                                                                                                                                                  APPLICANT: KIKUCHI, YASUfumi
TITLE OF INVENTION: APOPTOSIS-INDUCING SINGLE-CHAIN
FILE REFERENCE: 065678/0102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Norton, Elizabeth J. APPLICANT: Deikman, Alan B. TITLE OF INVENTION: Recombinant TITLE OF INVENTION: Antigen FILE REFERENCE: 00415-02
                                                                                                                           TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAGT 238
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                                                                                                                                                                                                                                                                                                                                            OH-EDA, Masayoshi
                                                                                                                                                                                                                                                                                                                                                                               FUKUSHIMA, Naoshi
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   Conservative
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                    69.9%;
70.7%;
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29;
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Score 900.5;
Pred. No. 3e-6
29; Mismatches
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Pred. No. 2e-69;
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                  5; DB 27;
3e-69;
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                                   Length 271;
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; SOFTWARE: Patentin
; SEQ ID NO 26
; LENGTH: 274
; TYPE: PRT
; ORGANISM: Mus sp:
US-09-523-095A-26
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                                                                                                                                                                                                                                                                                                                                                     Query Match
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APPLICANT: OH-EDA, Masayoshi
APPLICANT: KIKUCHI, Yasufumi
TITLE OF INVENTION: APOPTOSIS-INDUCING SINGLE-CHAIN
FILE REFERENCE: 065678/0102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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NUMBER OF SEQ ID NOS:
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                               238 TKLEIK 243
                                                               201 QSPKLLIYKVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYFCSQSTHVPYTSGGG
                                                                                              178
                                                                                                                            118
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261 TKLEIK 266
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                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                            1 QVQLQQSGAEI,VRSGASVKLSCTASGENIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
                                                                                           QSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAG 237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAG
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                                                                                                                                                         GGGSGGGGSGCGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRPG 177
                                                                                                                                                                                                                                                         QVQLQQSGPDLVKPGASVKMSCKASGYTFVNHVMHWVKQKPGQGLEWIGYIYPYNDGTKY 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QSPKLLIYKVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYFCSQSTHVPYTSGGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                      69.98;
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                                                                                                                                                                                                                                                                                                                         29; Mismatches
                                                                                                                                                                                                                                                                                                                                        Score 900.5; DE Pred. No. 3e-69;
                                                                                                                                                                                                                                                                                                                                                      DB 27; Length 274;
                                                                                                                                                                                                                                                                                                                       38;
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; Sequence 16, Application US/09364088

RESULT 10 US-09-364-088-16

GENERAL INFORMATION:

APPLICANT:

Yashwant

M. Deo, et al.

COMPOUNDS

ANTIBODIES

TITLE OF INVENTION: THERAPEUTIC COMPOUND TITLE OF INVENTION: OF ANTI-FC RECEPTOR NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:

ADDRESSEE:

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                                                                                US-09-523-279-16
                                                                                                       RESULT
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Sequence 16, Application US/09523279
GENERAL INFORMATION:
APPLICANT: Yashwant M. Deo et al.
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISES OF ANTI-FC RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 171;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDLIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: MXI-043CP2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 0 FILING DATE: 07-JUNE-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 07-JUNE-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
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                                                                                                                                                                 509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   278 IKLQQSGAELVRSGTSVKLSCTASGFNIKDSYMHWLRQGPEQGLEWIGWIDPENGDTEYA 337
                                                                                                                                                                                                    235 GAGTKLEIK 243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 553 amino
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                            62 PKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVTV 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (617)742-7414
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Pred. No. 7.7e-69;
7; Mismatches 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
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SEQ ID NO 16
LENGTH: 553
TYPE: PRT
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Best Local Similarity
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PRIOR FILING DATE: 1996-0
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1995-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1999-(PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-1
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CURRENT FILING DATE: 2000-03-10
PRICE APPLICATION NUMBER: 09/364,088
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                                                COUNTRY: USA:
ZIP: 02109-1375
COMPUTER READABLE FORM:
MEDIUM TYPE: .Floppy disk
.COMPUTER: IBM PC compatible
.COMPUTER: IBM PC compatible
.COMPUTER: PC-DOS/MS-DOS
.COMPUTER: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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CITY: Boston
STATE: Massac
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   175 RPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTF 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         338 PKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCNEGTPTGPYY---FDYWGQGTTVTV
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                WAPPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62 PKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVTV 114
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                                                                                                                                                                                            Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/09188082
                                                                                                                                                                                                                               60 State Street, Suite 510
                                                                                                                                                                                                                                                                                                                                                          Joel Goldstein
Robert Graziano
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VENTION: THERAPEUTIC COMPOUNDS COMPRISED
                                                                                                                                                                                                                                                                                                                                                                                              Yashwant M. Deo
                                                                                                                                                                                                                                                      LAHIVE & COCKFIELD
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Pred. No. 7.7e-69;
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; ORGANISM: Mus
US-09-523-095A-40
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                                                  Matches 173;
                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                 SEQ ID NO 40
LENGTH: 245
                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 40, Application US/09523095A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/523,095A
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: JP 11-63557
                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin V
                                                                                                                                                                                                                                             PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MX
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           395 SSGGGGSGGGGGGGSENVLTQSPAIMSASPGEKVTITCSASSSV-----SYMHWFQQ 448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     338 PKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCNEGTPTGPYY---FDYWGQGTTVTV 394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               278 IKLQQSGAELVRSGTSVKLSCTASGFNIKDSYMHWLRQGPEQGLEWIGWIDPENGDTEYA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62 PKFQGKATMTADTSSNTAYLQLSSLASEDTAYYYCN-----FYGDALDYWGQGTTVTV 114
Match 69.8%;
Local Similarity 68.7%;
les 171; Conservative 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RPGQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTF 234
                                                                                                                                                                                                                                                                                                                                                                               FUKUSHIMA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            553 amino acids
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                                                             Conservative
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70.3%;
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                                                        Score 892.5; DB 2/;
Pred. No. 1.3e-68;
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Pred. No. 7.7e-69;
7; Mismatches 35
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SEQ ID NO 34
LENGTH: 271
TYPE: PRT
Sequence 32, Application US/09523095A
GENERAL INFORMATION:
APPLICANT: FUKUSHIMA, Naoshi
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Best Local Similarity
Matches 173; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1999-
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CURRENT FILING DATE: 2000-03-10
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TITLE OF INVENTION: APOPTOSI
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                                                                                                        258 TKLEIK 263
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                                                                                                                                                                                               QSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAG
                                                                                                                                                                                                                                       GGGSGGGGSGVVMTQSPLSLPVSLGDQASISCRSSQSLVHSNGKTYLHWYLQKPG
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BER: JP 11-63557
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-APPLICANT:

OND,

Shinsuke

APPLICANT:

OH-ED&, Masayoshi

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SOFTWARE: PatentIn SEQ ID NO 32 LENGTH: 274 TYPE: PRT ORGANISM: Mus sp. US-09-523-095A-32
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Best Local Similarity 70.3
Matches 173; Conservative
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TITLE OF INVENTION: APOPTOSIS-INDUCING SINGLE-CHAIN FV
FILE REFERENCE: 065678/0102
CURRENT APPLICATION NUMBER: US/09/523,095A
CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: JP 11-63557
PRIOR FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 40
SOFTWARPS: DELICATION SEQ ID NOS: 40
                   238 TKLEIK 243
                                                                                                                                     261 TKLEIK 266
                                                                                      178 QSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAG 237
                                                                    201 QSPKLLIYKVSNRFSGVPDRFSGSGSVTDFTLMISRVEAEDLGVYFCSQSTHVPYTFGGG 260
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US-08-661-052-16

3 US-09-184-658-3

1 US-08-331-3978-34

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2 US-08-375-811-53

3 US-08-875-811-53

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3 US-08-8279-728-8

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1 US-08-461-880-4

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- SS - GT - GT - GT - GT - GT - GT - GT	OB-652-507-2 equence 2, Application US/086; atent No. 5876691 atent No. 5876691 TITLE OF INVENTION: APPLICANT: INVENTION: APPLICANT: INVENTION: NUMBER OF SEQUENCES: CORRESSONDENCE ADDRESS: ADDRESSE: Nixon 6 Vande; STREET: 1100, No. 5876691; CITY: Arlington STATE: V1rginia COUNTR: USA ZIP: 2201-4714 COMPUTER READABLE FORM: MEDIUM TYDE: Floppy disk COMPUTER: IBM PC compatil OPERATING SYSTEM: PC-DOS, SOSTWARE: Patentin Relea; CURRENT APPLICATION DATA: APPLICATION NUMBER: US/00) FILING DATE: 02-Jul-1996 CLASSIFICATION. 530 ATTORNEY/AGENT INFORMATION: NAME: Arthur R. Crawford REGISTRATION NUMBER: 25: APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: ATTORNEY/AGENT INFORMATION PREFERENCE DOCKET NUMBER: 25: APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: ATTORNEY APPLICATION ACIDS REFERENCE TOWBER: 25: APPLICATION TOWBER: 25: ATTORNEY APPLICATION TOWBER: 25: APPLICATION TOWBER: 25: ATTORNEY APPLICATION TOWBER: 25: ATTORNEY APPLICATION TOWBER: 25: ATTORNEY APPLICATION TOWBER: 25: ATTORNEY APPLICATION TOWBER: 25: ATTORNEY APPLICATION TOWBER: 25: APPLICATIO	00000000000000000000000000000000000000
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QKPGTSPKLWIYSTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPLT

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VSSGGGGSGGGGGGGGDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLL 173 APKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCNEGTPTGPYY---FDYWGQGTTVT

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US-08-661-052-16
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Best Local Similarity
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                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino acid
                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: MX TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 07-JUNE-199
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
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338 PKFQGKATFTTDTSSNTAYLQLSSLTSEDTAVYYCNEGTPTGPYY---FDYWGQGTTVTV 394
                                                                278 IKLQQSGAELVRSGTSVKLSCTASGFNIKDSYMHWLRQGPEQGLEWIGWIDPENGDTEYA 337
                  S2 PKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN-----FYGDALDYWGQGTTVTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Arnold, Beth E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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NVENTION: THERAPEUTIC COMPOUNDS COMPRISED
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 63
LENGTH: 289
                                                           Best Local Similarity Matches 167; Conserv
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Best Local s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/184,658
CURRENT FILING DATE: 1998-11-02
EARLIER APPLICATION NUMBER: 60/065,423
EARLIER FILING DATE: 1997-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: SIGNAL
LOCATION: (1)..(22)
OTHER INFORMATION: I
OTHER INFORMATION: 1
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                                                                                                                                                      OTHER INFORMATION:
                                                                                                                                                                   NAME/KEY: SITE
LOCATION: (284).
                                                                                                                                                                                                 LOCATION: (270)..(;
OTHER INFORMATION:
FEATURE:
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LOCATION: (139)..(154)
OTHER INFORMATION: 16
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LOCATION: (23)...
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                                                                                                                                                                                                                                              NAME/KEY: SITE
                                                                                                                                                                                                                                                                              OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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LOCATION: (155).
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23 EVQLVESGGGSVQPGGSLKLSCAASGFTFNTYGMSWVRQTPDKRLEWVATINSNGGLTFY 82
              1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
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DRMATION: 5109
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Pred. No. 7.9e-61;
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US-08-331-398A-34
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                                       Query Match
Best Local Similarity
                                                                                                                                                                                                 REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 015280-126110US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-OCT-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Single Chain B3 Antibody Fusion Proteins TITLE OF INVENTION: and Their Uses (as amended) NUMBER OF SEQUENCES: 68
                                                                                                                                                                                                                                                                                                                                           FILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
NAME: Hunter, Tom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                           LENGTH:
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One Market Plaza, Steuart Street Plaza
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FitzGerald, David
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                                        64.3%; Score 828.5; 63.7%; Pred. No. 1.:
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                       34;
                       Mismatches
                                        .1e-59;
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                       Indels
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 Query Match 64.3
Best Local Similarity 63.7
Matches 156; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: !Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver:
CURRENT APPLICATION DATA:
'APPLICATION WIMBER: US/08/331,397B
FILING DATE: '28-CCT-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                          TELEFAX: (415) 543-50 INFORMATION FOR SEQ ID NO:
                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 248 amino acids
                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                                                                                       FILING DATE: 12-OCT-1990 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
TITLE OF INVENTION:
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APPLICATION NUMBER: US 07/767,331
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ADDRESSEE: Townsend
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                                                                                                                                                                                                                                                                                              NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
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One Market Plaza, Steuart Street
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                 64.3%;
63.7%;
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Specific Antibody Fragments, Fusion Proteins,
   34;
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Score 828.5; DB 2;
Pred. No. 1.1e-59;
4; Mismatches 52;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/331,398
FILING DATE: 28-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
PRIOR APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-OCT-1990
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APPLICATION NUMBER: US,
FILING DATE: 03-DEC-19
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MEDIUM TYPE: Floppy disk
                                                                            SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: Fusion Proteins, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David
TOPOLOGY: 1
                                                                                                                                          REFERENCE/DOCKET NUMBER: 015280-126140US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                        LENGTH:
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                                                                                                                                                                                                         NAME: Weber, Ellen L. REGISTRATION NUMBER: 32,762
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                                   amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Firzuerald, David
Brinkmann, Ulrich
                                                                                                                        (415) 576-0300
                   linear
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Best Local Similarity
Matches 156; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 53, Application US/08875811 Patent No. 6045793
               TELEFAX: (415) 576-030
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 365 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                          REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION UNMBER: US 60/011,800
FILING DATE: 21-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend
                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/OFFILING DATE: 19-FEB-1998
                                                                                                                                            NAME: Faris, Susan K. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  63 DTVKGRFTISHDNARNTLYLQMSRLKSEDTAIYSCARGLAWGAWFAYWGQGTLVTVSSGG 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62 PKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCN---FYGDALDYWGQGTTVTVSSGG
                                                                                     TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SPKLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYYCFQGSHVPFTFGSGT
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amino acid
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               365 amino acids
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Newton, Dianne L.
Boque, Lluis
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VENTION: Recombinant Ribonuclease Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                            (415)
                                                                        576-0300
                                                                                         576-0200
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                                                                                                                                                                                                                                                                                                                                                                                     Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                   US/08/875,811
                                                                                                                                              41,739
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                                                                                                                               015280-244100US
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US-08-875-811-55
; Sequence 55, App
; Patent No. 60457
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Best Local Similarity 61.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 150;
                                                                           ATTORNEY/AGENT INFORMATION:
NAME: FAI'S, SUSAN K.
REGISTRATION NUMBER: 41,739
REFERENCE/DOCKET NUMBER: 0152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
               TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 55: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tent No.
                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Wlodawer, TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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  ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               San Francisco
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366 amino acids
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Newton, Dianne L.
Boque, Lluis
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US-08-279-772A-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 63.0
Best Local Similarity 61.7
Matches 150; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Russel
APPLICANT: Fuller
                                                                                TELEPHONE:
TELEFAX: 60
INFORMATION FOR
                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27,386
REFERENCE/DOCXET NUMBER: 11-7
TELECOMMUNICATION INFORMATION:
-: TELEPHONE: 638-251-5000
                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/279,772A
                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MOLECULE TYPE:
                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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CITY: Madison
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              TYPE: amino TOPOLOGY: li
                                                                                                                                                                                                                      CLASSIFICATION: 435
                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: United States of America ZIP: 53701-2113
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                                              LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                               ΨH
                                                239 amino acids
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                                                                                                  608'-251-9166
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Fuller, James
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                                 acid
                                                                                  ID NO:
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Cells
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; MOLECULE TYPE: US-08-902-486-11
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Best Local
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                                                                                                                                                                                                                                                                   TELEFAX: 608-25
                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 67
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   177 KLWIYTTSNLASGVPARFSGSGSGTSYSLTISRMEAEDAATTYCHQRSTYPLTFGSGTKL 236
                                                                                                                                                                                                                             LENGTH: 239 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  241 EIK 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity hes 151; Conserv
                                                                                      Local Similarity
                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYCNFYGDALDYWGQGTTVTVSSGGGG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.
QVQLQQSGAELAKPGASVKMSCKASGYTFTSYRMHWVKQRPGQGLEWIGYINPSTGYTEY 62
                                   QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ELK
K
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                                                                                                                                                                                                                                                                                      608-251-9166
                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Quarles & Brady
                                                                                                                                                                                                                                                                                                        608-251-5000
                                                                                                                                                                             protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    James
                                                                                        62.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 METHOD FOR PRODUCING ANTIBODIES AND PROTEIN TOXINS IN PLANT CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                 US/08/902,486
                                                                                                                                                                                                                                                                                                                                          670513.90261
                                                                    Score 804; DB 3;
Pred. No. 9.6e-58;
4; Mismatches 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 9.6e-58;
4; Mismatches 52;
                                                                                                       Length 239;
                                                                      Indels
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US-08-463-163-3
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                                                                                                                                                         REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 22-JAN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
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                                                                                                                                                                                                              NAME: Weber, Ellen L. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 05-JUI
                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
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                                   ECULE TYPE: protein
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                                                                  amino
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                                                                                    599 amino acids
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                                                                                                                                          (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FitzGerald, David J. Chaudhary, Vijay K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Queen, Cary L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pastan,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Townsend and Townsend and Crew e Market Plaza, Steuart Street Tower
                                                                                                                                                                                                                                                                                                                                                                           UMBER: US 06/911,227
                                                                                                                                                                                                                                                                     08-APR-1992
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21-APR-1989
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Matches

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; MOLECULE TYPE:
US-08-752-844-66
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US-08-752-844-66
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 Query Match
Best Local Similarity
Matches 154; Conserv
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Best Local Similarity
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                                                                                                                                                                                TELEFAX: (415) 494-079
TELEX: 706141
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                               REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 30.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Foon, Kenneth A. APPLICANT: Chatterjee, Sunil K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: Schiff, J. Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        181 KRLIYLVSKLDSGVPDRFTGSGSGTDETLKINRVEAEDLGVYYCWQGTHSPLTFGAGTKL 240
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                                                                                                                                TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    62 NQKFKDKATLTADKSSSTAYMQLSSLTFEDSAVYYCARGGGVFDYWGQGTTLTVSSGGGG 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                              263 amino acids
                                                                                                                                                                                                                 (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chatterjee, Malaya
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                                                                                                                   linear
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                  62.2%;
62.3%;
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   36;
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Score 801.5; I
Pred. No. 1.7e
36; Mismatches
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Pred. No. 2.8e-57;
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                 .7e-57
                                  DB 2;
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US-08-447-402-1
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                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                              REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (512) 474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM.PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 260 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                              CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              CLÁSSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Chen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      197
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                                                                                                                        TELEFAX: 19-092
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                                                                                                                                                                                                                                                                             FILING DATE: 10-JUN-1994
CLASSIFICATION: 424
APPLICATION NUMBER: 07/7
FILING DATE: 15-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA .
ZIP: 77210-4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: P. O.,
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TLE OF INVENTION:
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                                                TOPOLOGY:
                                                                                                                                                                                                              NAME: Kitchell, Barbara S. REGISTRATION NUMBER: 33,928
                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GTKLEIK 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GQSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYYCWQGTHSPLTFGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGGGSGGGSGGGSDVLMTQTPLTLSVTIGQPASISCKSSQSLLDSDGKTYLNWLLQRP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QVQVKESGPFLVPPSQSLSITCTVSGFSLTTYGVSWIRQPPGKGLEWLGAIWGD---GTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GQSPNLLIYFVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYYCFQGSHVPWTFGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QVQLQQSGAELYRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIG--WIDPENGDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                              TYPE: protein
                                                               amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08447402
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                                                                                260 amino acids
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Georgiou, George
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FR: 07/794,731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                               NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tent No.
                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS DOS SOTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                             FILING DATE: 27-FEB-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: OPPERMANN, HERMANN APPLICANT: TIMASHEFF, SERGE N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             238 TKLEIK 243
                                                                                                                                             NAME: CAMPBELL ESQ, PAULA A REGISTRATION NUMBER: 32,503 REFERENCE/DOCKET NUMBER: CRI
                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER:
                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION: 435
                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                ELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGGSGGGGSGGGSDIVLTQSPASLAVSLGQRATISCRSSQSLVHSNGNTYLNWYQQKPG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NQKFKGKATLTVDKSSSTAYMELRSLTSEDSAVYYCAGSSGNKWAMDYWGHGASVTVSSG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                01748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HOPKINTON
                                               amino acid
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                                                               246 amino acids
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                             linear
                                                                                                             617/248-7000
                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
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Query Match:

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                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/133,804
                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Kelley; Robin D.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                     MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                            LENGTH: 252 amin TYPE: amino acid
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CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 QVQLQQSGAELVRSGASVKLSCTASGFNIKDYYMHWVKQRPEQGLEWIGWIDPENGDTEY
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          35,
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   Score 795.5; DB 1
Pred. No. 4.9e-57;
5; Mismatches 49
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; Mismatches
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Searc Job t	Дb	Qy	DЪ	Qy	Db	Qy	Дb	Qy	DЪ
th con	240	238	180	178	121	118	62	61	N
Search completed: February 12, 2001, 15:23:45 Job time: 59 sec	TKLEIK 245	TKLEIK 243	QSPKLLIYKVSNRFSGVPDRFSGSGSGTDFTLKISKVEAEDLGIYFCSQTTHVPPTFGGG 239	QSPKRLIYLVSKLDSGVPDRFTGSGSGTDFTLKINRVEAEDLGVYYCWQGTHSPLTFGAG 237	- SGSSSGSSSSGSDVVMTQTPLSLPVSLGDQASISCRSSQSLVHSNGNTYLNWYLQKAG 179	GGGSGG	62 NQKFKGKATLTVDKSSSTAYMELRSLTSEDSAVYYCAGSSGNKWAMDYWGHGASVTVSS- 120	APKFQGKATMTADTSSNTAYLQLSSLASEDTAVYYC-NFYGDALDYWGQGTTVTVSSG 117	:

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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gb_est55:* gb_est54:* gb_est51:*

gb_est49: gb_est48:*

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gb_est62

em_estro em_estro16:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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RESULT 1
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A3505978 IN0346U72

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A300651 EST13764

BE307170 601098164

AW407858 UI-HF-BL0

BE3287568 601097304

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AZ447391 IN0244H01

BF022578 uy51a07.y

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AW403810 UI-HF-BK0

AW402541 UI-HF-BK0

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BF016722 uy34h12 y
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AW390292 CM2-ST018
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AW404610 UI-HF-BLO
BE281961 601099518
AW951579 EST363649
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AW406572 UI-HF-BL0
AW406512 UI-HF-BL0
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ACCESSION VERSION KEYWORDS SOURCE BE306420. 1057 bp mRNA EST 13-JUL-2000 601103347F1 NCI_CGAP_Lu29 Mus musculus cDNA clone IMAGE:3495755 5', mRNA sequence. BE306420 BE306420.1 GI:9159604 EST. house mouse.

em_gss16:* em_gss17:* em_gss7:*
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Plate: LLAM8546 row: n column:
High quality sequence stop: 695.
Location/Qualifiers
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CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              National Institutes of Health, Mammalian Gene Collection (MGC)
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Clone distribution: MGC clone distribution information can
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                                                                                       434
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Organ: lung; Vector: pcMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo: Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH" a 289 c 273 g 221 t
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/clone_lib="NCI_CGAP_Lu29"
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TATTATAGTAGTCCTCGGACGTTCGGCCAGGGGACCAAGGTGGAAATCAAA 391
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Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
CDNA Library Preparation: M.B. Soares Lab
CDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information
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Tel: (301) 496-1550
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Constructed from size fractionated cytoplasmic mRNA (1.5-2.5kb). Directionally cloned. Cells provided by Low M. Staudt, Ph.D. Library preparation by Maria de Fatima Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
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/tissue_type="lymph"
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                     catccactagggaatctggagtccctgatcgcttcacaggcagtggatctgggacagatt 640
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CATCTACCCGGGAATCCGGGGTCCCTGACCGATTCAGTGGCAGCGGGTCTGGGGACAGATT
                                                                                      ACTACTTAGCTTGGTACCAGCAGAAACCAGGACAGCCTCCTAAGCTGCTCATTTACTGGG
                                                                                                                                                                            AGAGGGCCACCATCAACTGCAAGTCCAGCCAGAGTGTTTTATACAGCTCCAACAATAAGA 123
                                                                                                                                                                                                                                                                     GTCGTGACGGGGACATCGTGATGACCCAGTCTCCAGACTCCCTGGCTGTGTCTCTGGGCG 63
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IL5-UM0071-120400-065-d06
AW802126
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This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL
Project. This entry can be seen in the following URL
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400-065-d06&t3-2000-04-12&t4=1)
Seq primer: puc 18 forward
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
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1. (bases 1 to 533)
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NIH-MGC http://www.ncbi.nlm.nlh.gov/MGC/.
National Institutes of Health, Mammalian Gene Collection
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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//note="Vector: pT/T3-Pac; Site_1: Not1; Site_2: Eco RI;
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/tissue_type="lymph"
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/cell_line="MGC85"
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ccgaaagaattacttggcttggtatcagcagaaaccagggcagtctcctaaagtgctgat 572
                                                                                                   tottataatotacog---acgttcggcgggggcaccaagctggaaatcaaa 747
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Seq primer: M13 Forward.
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Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, Mammalian
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/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
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/tissue_type="lymph"
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                  ggtggcggatcgga&attgagctcacccagtctccatcttccctggctgtgtcagcagga 459
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TCAGCAATATTATAGTATTCCCTTCACCTTCGGCCAAGGGACACGACTGGAGATTAAA 358
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267; Conserv
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Unpublished (1999)
Contact: Simpson A.J.G.
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Seq primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=CM2&t2=CM2-ST0182-
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Ludwig Institute for Cancer Research
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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FAPESP/LICR Human Cancer Genome Project
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                                                                                                                                                                                                                                                                                            /note "Organ: stomach; Vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of
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173 c 162 g 146 t
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/clone_lib="ST0182"
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Contact: Robert St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     uw89h05.y1 Soares_mammary_gland_NMLMG Mus musculus cDNA clone IMAGE:3469209 5' similar to SW:HV07_MOUSE P01751 IG HEAVY CHAIN V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; 1 (bases 1 to 358)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           act: Robert Strausberg, Ph.D.
(301) 496-1550
                                                                                                                 Conservative
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                                                                                                                                                                                                                     /note-"Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from mammary gland tissue from a lactating female, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. " 93 g 79 t
                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="mammary gland"
/lab_host="DH10B"
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/sex="female (lactating)"
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/db_xref="taxon:10090"
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74.7%;
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"" ematches 86;
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                                                                                                                                                                                                                                                                                                                                   Eco RI site shown at the beginning of the se
Tissue Procurement: Louis M. Staudt, M.D.,
CDNA Library Preparation: M.B. Soares Lab
CDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AW404610 TSO bp mRNA UI-HF-BL0-acc-a-06-0-UI.rl NIH_MGC_37 IMAGE:3058451 5', mRNA sequence.
                                                                                                                                                                                                                                                                                    Seq
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Contact: Robert St
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National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                                                                                                                         : Robert
                                                                                                                                                                                                                                                                                                                     through the I.M.A.G.E. Consortium/LLNL
            /lab_host="DH108 (LTI)"
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
/constructed from size fractionated cytoplasmic mRNA
(1.5-2.5kb). Directionally cloned. Cells provided by Lo
M. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

150 c 139 g 121 t
                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3058451"
                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                   /cell_type="germinal center
/cell_line="MGC85"
                                                                                                                                                                    /clone_lib="NIH_MGC_37"
/tissue_type="lymph"
                                                                                                                                                                                                                                                                                                    .gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                         _Strausberg@nih.gov
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 644)
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601099518F1 NCI_CGAP_Lu29 Mus
mRNA sequence.
BE281961
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Plate: LLAM8537 row: c column:
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.
National Institutes of Health, Mammalian Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue Procurement: Gilbert Smith, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999
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                                 Site_2: NotI; Cloned unidirectionally. PLIDTARY constructed by Life Technologies. providing samples: Gilbert Smith, NIH* a 176 c 155 g 166 t
                                                                                  The rest Thi0B"

/lab_host="DHi0B"

/note="Organ: lung; Vector: pCMV-SPORT6; Site_1: sa.r.,
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/note="Organ: lung; Vector: pCMV-SPORT6; Site_1: sa.r.,
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/clone_lib="NCI_CGAP_Lu29"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Mus musculus"
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  264;
                                                                                                                                                                                                        Unpublished (2000)
Contact: John Quackenbush
The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
Tel: 301 838 3528
Fax: 301.838 0208
                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 388)
Hegde,P., Qi,R., Abernathy,K.,
,I.E., Saeed,A.I., Sharov,V., I
                                                                                                                                                                                                                                                                                                                                                                                                                                               AW951579: 388 bp mRNA EST 01-JUN-2000 EST363649 MAGE resequences, MAGB Homo sapiens cDNA, mRNA sequence. AW951579
                                                                                                                                                                     Seq primer: Reverse.
                                                                                                                                                                              Email: johnq@tigr.org
                                                                                                                                                                                                                                                                                      Assessment of gene expression patterns metastasis using a 19,200 element cDNA
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                         .//organism="Homo sapiens"
//db_xref="taxon:9606"
./clone_lib="MAGE resequences, |
//note="Vector: pBluescriptSKm"
a 110 c 97 g 94 t
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73.5%;
             27.0%;
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Sharov,V., Lee,N.H., Yeatman,T.J.
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  0,:
             Score 201.8; DB 9
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Seq primer:
High quality
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                                                                                                                                                                                                                                                                                                CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNI) DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov/image/html/iresources.shtml
                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                         Tissue Procurement: Gilbert Smith, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                              Email:
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          /note-"Organ: lung; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: Sall; transgenic model WNT-1, expression driven by MMTV-LTR enhancer; Cloned uniformally. Primer: Oligo dT. Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH" 138 c 126 g 120 t
                                                                                                    /strain="CZECH II"
/db_xref="taxon:10090"
/clone="IMAGE:3658670"
/clone_lib="NCI_CGAP_Lu30"
/tissue_type="tumor, metastatic to mammary"
/lab_host="DH10B"
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/strain="CZECH II"
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                                                                                                                                                                                                                                          Email: Robert_Strausberg@nih.gov
Eco RI site shown at the beginning of the sequence.
Eco RI site shown at the beginning of the sequence.
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 328)

NIH-MCC http://www.ncbi.nlm.nih.gov/MGC/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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UI-HF-BL0-acr-f-09-0-UI.rl NIH_M
IMAGE:3060232 5', mRNA sequence.
                                                                                                                                                                                              www-bio.llnl.gov/bbrp/image/image.html
Seq primer: M13 Forward.
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                                                                                                                                                                                                                                                                                                                                                                                                                             (301) 496-1550
                                       /db_xref="taxon:9606"
/clone="IMAGE:3060232"
/clone_11b="NIH_MGC_37"
/tissue_type="1ymph"
/cell_type="germinal center
/cell_line="MGC85"
                                                                                                          /organism="Homo sapiens"
/db_xref="təvo"
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                                                                                                                                                                                                                                                                                                    Tissue Procurement: Gilbert Smith, Ph.D.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus (bases 1 to 464)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence.
BE368025
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601221914F1 NCI_CGAP_Lu29 Mu
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                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
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Clone distribution: MGC clone distribution information can
                                                                                                                                                                                               quality sequence stop: 462.
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                                                                                                                                                                                                                                                                                                                                                           Robert_Strausberg@nih.gov
                                                                 /organism="Mus musculus"
/strain="CZECH II (feral)"
/db_xref="taxon:10090"
/clone="IMAGE:3590907"
/clone_lib="NCI_CGAP_Lu29"
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84 g 74 t
                  /tissue_type="spontaneous
Stem cell origin."
/lab_host="DH10B"
                                                                                                                                                                            ocation/Qualifiers
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79.5%;
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itutes of Health, Mammalian Gene Collection (MGC)
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             MGI:1422279
Seq primer:
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uy34h12.y1 NCI_CGAP_Lu30 Mus mus
similar to SW:HV02_MOUSE P01746
                                                          CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
image.llpl.gov/image/html/iresources.shtml
                                                                                                                                                        Email: Robert_Strausberg@nih.gov
Tissue Procurement: Gilbert_Smith, ph.D.
                                                                                                                                                                                        Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg,
Tel: (301) 496-1550
                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National_Cancer Institute, Cancer Genome Anatomy
                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammmalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1_ (bases 1 to 406)
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BF016722.1
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primer: -40RP from Gibco.
Location/Qualifiers
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Pred. No. 3.8e-48;
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IMAGE:3660143
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                                                                                                          This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be rein in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM4&t2=PM4-HT0348-261199-001-B11&t3-1999-11-26&t4=1)
                                                                                                                                                                                                                                                                                      Contact: Simpson A.J.G.

Laboratory of Cancer Genetics

Ludwig Institute for Cancer Research

Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                       Seg primer: puc 18 forward
High quality sequence start: 10
High quality sequence stop: 441
                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AW383565 459 bp mRNA EST 04-FEB-2000 pM4-HT0348-261199-001-B11 HT0348 Homo sapiens cDNA, mRNA sequence. AW383565 AW383565.1 GI:6888133
                                                                                                                                                                                                Email: asimpson@ludwig.org.br
                                                                                                                                                                                                                                                                      Brazi
                                                                                                                                                                                                                                                                                                                                                                               HCGP http://www.ludwig.org.br/ORESTES.
The FAPESP/LICR Human Cancer Genome Project.
Unpublished (1999)
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                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="tumor, metastatic to mammary"
/lab_host="DH10B"
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/clone="IMAGE:3660143"
/clone_1ib="NCI_CGAP_Lu30"
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Pred. No. 1.6e-47;
0; Mismatches 57; I
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Best Local Similarity
Matches 245; Conserv
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                                                                                                                      ctggaaatcaaa 747
                                                                                                                                                                             gcagtttattactgcaagcaatcttataatctaccg---acgttcggcggggggcaccaag 735
                                                                                                                                                                                                                               GGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGGCTGAAGATGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note-"Organ: head_neck; vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196.716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA maplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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/dev_stage="Adult"
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78.5%;
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Pred. No. 1.5e-46;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: sp_bacteria
3: sp_fungi:*
4: sp_human:*
5: sp_inverteb:
6: sp_mammal:*
7: sp_mac:*
8: sp_organell:
9: sp_bage:*
10: sp_plant:*
11: sp_rodent:
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13: sp_vertebr
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Q9u183 homo
Q9u194 homo
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Q9j177 mus
Q9qxf0 mus
Q9qxf0 mus
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Q91185 mus
Q921c4 mus
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Q9qxe9 mus
Q9u181 homo
Q9u192 homo
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Q9N0W4	Q9N0W6	Q9UL72	Q9UL91	095973	Q9N0W5	095978	Q9JL76	Q9UL71	Q9UL73	Q9UL93	Q9UL88	Q9U410	Q9UL90	Q9Y509	Q9JL83	Q9UL89	Q9JL84	Q9JL80	Q9JL78	Q9UL80	Q9R1A5	Q9Z1C6	Q9UL86	Q9JL82	Q9Y298
Q9n0w4 oryctolagus	Q9n0w6 oryctolagus	Q9u172 homo sapien	Q9u191 homo sapien	O95973 homo sapien	Q9n0w5 oryctolagus	095978 homo sapien		Q9ul71 homo sapien	hoπo	homo	Q9u188 homo sapien	Q9u410 schistosoma	homo	Q9y509 homo sapien	3 mus	Q9ul89 homo sapien	Bum	Q9j180 mus musculu		homo	Q9r1a5 mus musculu	mus	-	2 mus	Q9y298 homo sapien

ALIGNMENTS

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Qy 1 QVKLQESGAEĹVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY 60 : : : : : : : : :	Query Match 56.6%; Score 744.5; DB 11; Length 298; Best Local Similarity 56.4%; Pred. No. 7.4e-58; Matches 141; Conservative 37; Mismatches 61; Indels 11; Gaps 3;	SQ SEQUENCE 298 AA; 31867 MW; E0F96B8A17004317 CRC64;		Submitted (DEC-1999) to the		-	RA Shinohara N. Demura T. Fukuda H.:	•	Submitted (DEC-1999)			NCBI_TaxID=10090;	OC Eukaryota; metazoa; cuoruata; craniata; vertebrata; Euteteoscomi; OC Mammalia; Eutheria; Rodentia; Sciurconathi; Muridae; Muripae; Mus.	Mus musculus (Mouse).	CZ	01-OCT-2000 (TrEMBLrel. 15,	01-MAY-2000 (TramBirel. 13,	01-MAY-2000 (TrEMBLrel.	Q9QYEO;	ID OGOVEO DREITMINARY: DRT: 298 AA.	DQQYFQ F	Depart m 1
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O1-OCT-2000 (TrEMBLrel. 15, Last annotation update)
O1-OCT-2000 (TrEMBLrel. 15, Last annotation update)
ANTI-MYOSIN IMMUNOGLOBULIN HEAVY CHAIN VARIABLE REG
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
"""""" ia: Eutheria; Rodentia; Sciurognathi; Murida
                                                                                                                                                                                                                                                                                                                                        01-MAY-1999 (TIEMBLIEL 10, Created)
01-MAY-1999 (TIEMBLIEL 10, Last sequence update)
01-MAY-2000 (TIEMBLIEL 13, Last annotation update)
ANTI-PORCINE VCAM MAB 3F4 HEAVY CHAIN VARIABLE REGION
MUS MUSCULUS (MOUSE).
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SEQUENCE
                                   STRAILE J.P., Glannoni M.A., Hart
Musiler J.P., Glannoni M.A., Hart
Matis L.M., Evans M.J.;
"Humanized porcine VCAM-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Malkiel S., Liao L., Cunningham M.W., Diamond B.; "Characterization of cross-reactive monoclonal anti-myosin/anti-n-acetyl-glucosamine antibodies from mice with autoimmune myocarditis."; Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.

EMBL: AF206021: AAF69319.1; -.
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEYAPNFQGKA 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TPSLKDKFIISRDNAKNTLYLQMSKVRSEDTALYYCARASYYGHSA----YWGQGTTVTV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA;
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11944 MW;
                                                                                                                                                                                                                                                                                   Chordata;
Rodentia;
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Pred. No. 4.9e-31,
1; Mismatches 1;
                                                                                                              Hartman
                                                                                                                                                                                                                                                                                   Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
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                                          monoclonal
                                                                                                       S.L.,
       leukocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  118
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                                                                                                       Elliott E.A.,
   antibodies
yte binding
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                                                                                                                                                                                                                                                                                                                                                                                          (FRAGMENT).
to porcine
                                          with chimeric
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Clemens A., Rademaekers A., S
Submitted (DEC-1997) to the E
EMBL; AJZ25174; CAB65237.1; -
INTERPRO; IPR003006; .
PFAM; PF00047; ig; 1.
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Q9QXE9;
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SEQUENCE
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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Submitted (NOV-1996) to tl
EMBL; U78801; AAD00293.1;
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                                                                                                                                                                        SS 122
                                                                                                                                                                                                                                                                                                               QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
:|:||:|| ||| :|| :|| :|| :| | | |
EVQLQQSGPELVKPGASVKMSCKASGYTFTDYYMKWVKQSHGKSLEWIGDINPNNGGTSY
                                                                                                                                                                                                                                                         APNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTV 120
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                                                                                                                                                                                                                                NQKFKGKATLTVDKSSSTAYMQLNSLTSEDSAVYYCARDRY-----YAMDYWGQGTSVTV
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118
118 AA;
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13036 MW;
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Rođentia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MW;
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, Last annotation update)
V-D-J REGION (FRAGMENT).
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Pred. No. 1.2e
13; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 379.5; DB
Pred. No. 3e-26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDDE2AF84D499734
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No. 1.
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; Murinae; Mus
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01-JUN-2000 (TIEMBLIEL 13, Last sequence update)
01-JUN-2000 (TIEMBLIEL 14, Last annotation update)
MYOSIN-REACTIVE IMMUNOSLOBULIN HEAVY CHAIN VARTHER
Eukaryota; Metaron
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MEDLINE-98277139; PubMed-9614934;
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Eukaryota; Metazoa; Chordata; Craniata; V
Mammalia; Eutheria; Primates; Catarrhini;
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                QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
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 EVQLVESGAEVKKPGASVKVSCKASGYTFSSYYMHWVRQAPGQGLEWMGIINPSGGSTSY
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Primates;
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                          EMBL; AF
                                                Malkiel S., Liab L., Cunningham M.W., Diamond B.; "Characterization of cross-reactive monoclonal anti-myosin/anti-n-acetyl-glucosamine antibodies from mice with autoimmune myocarditi Submitted (NOV:1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                   Eukaryota; Metażoa;
Mammalia; Eutheria;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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(TrEMBLrel. 13,
(TrEMBLrel. 15,
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Rodentia;
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Pred. No. 1
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Best Local S
Matches 72
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01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation updat
ANTI-MYOSIN IMMUNOGLOBULIN HEAVY CHAIN VARIABLE F
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
    STRAIN-DBA/2
                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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MEDLINE-98277139; PubMed-9614934;
Wu X., Liu B., Van der Merwe P.L.,
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Mammalia; Eutheria;
NCBI_TaxID=9606;
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                     SEQUENCE FROM N.A
                                                              NGBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 368.5;
Pred. No. 2.
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Pred. No. 2.3e-25;
                                                                                 Sciurognathi;
                                                                                                       Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                     PRT;
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EMBL; AF206029; AAF69327.1; -
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae
MCBI_TaxID=10090;
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                        , Last sequence update)
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Pred. No. 3.
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"Characterization of cross-reactive monoclonal anti-myosin/anti-n-
acetyl-glucosamine antibodies from mice with autoimmune myocarditis.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, AF206025; AAF69323.1;
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INTERPRO;
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MEDLINE=98277139; PubMed=9614934;

Van der Merwe P.L.,
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01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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01-MAY-2000
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Mammalia; Eutheria;
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QSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTSWTFGEGTKVEIK 107
                         ESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLP-TFGGGTKLEIK
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                                                                                                                                                                                                                                                                                                                                                         ig; 1.
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11738 MW;
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Rodentia;
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Primates;
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Pred. No. 3.7e-25;
                                                                                                                                                                          Score 365.5; DB 4;
Pred. No. 4.6e-25;
1; Mismatches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                87:184-192(1998)
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Catarrhini;
                                                                                                                                                                                                                                                                                          C06681716C4D16F3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              404885FDE6BA56F8 CRC64;
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1; Hominidae;
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; Homo.
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Q9UL79;
01-MAY-2000
                                                                                                                                                                                                                                                            Malkiel S., Liao L., Cunningham M.W., Diamond B.; "Characterization of cross-reactive monoclonal anti-myosin/anti-n-acetyl-glucosamine antibodies from mice with autoimmune myocarditi. Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases. EMBL; AF206031; AAF69329.1; -.
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01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
ANTI-MYOSIN IMMONOGLOBULIN HEAVY CHAIN VARIABLE REGION (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Mammalia; Eutheria; NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF035035; AAI
HSSP; P01607; 1REI
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (Human)
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                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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                           ELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEYAPNFQGKAT 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QSGVPSRFSGSGSGTDFTLTISCLQSEDFATYYCQQYYSFPPTFGQGTKVEIK
ELVKPGASVKÁSCKASGYTFTSYVMHWVKQKPGQGLEWIGYINPYNDGTKYNEKFKGKAT
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                                                                                      Similarity 59.1
70; Conservative
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Rodentia;
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Pred. No. 7e-25;
9; Mismatches
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                                                                                      Score 359; DB 11;
Pred. No. 1.8e-24;
5; Mismatches 19
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GenCore version 4.5
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OM protein - protein search, using sw model

Run on: February 12, 2001, 15:25:01; Search time 26.51 Seconds (without alignments) 637.769 Million cell updates/sec

Title: US-09-297-181-4
Perfect score: 1315
Sequence: 1 QVKLQESGAELVRSGASVNL......YCKQSYNLPTFGGGTKLEIK 249

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5
Searched: 195891 seqs, 67900655 residues

Total number of hits satisfying chosen parameters: 195891
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

PIR_66:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

45	44	43	42	41	40	39	38	37	36	35 5	34	ω ü	32	31	30
463.5	464	465	466	467	467.5	467.5	468	468.5	468.5	469	470.5	474.5	474.5	475	477.5
35.2	35.3	35.4	35.4	35.5	35.6	35.6	35.6	35.5	35.6	35.7	35.8	36.1	. 36 . 1.	36.1	36.3
113	133	97	112	221	118	113	116	112	112	122	240	120	113	138	214
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S30520	K4HUJI	A42575	S09970	S49220	S25174	JC2270	S15672	F30538	E30538	S06823	S06084	S03471	PL0264	S26040	S68212
Ig kapp	Ig kapp	Ig kapp	Ig kapp	Ig gamma-	Ig heavy	PL7-6 an	Ig heav	Ig kapp	Ig kapp	Ig heavy	. Ig kapp	Ig heav	Ig kapp	. Ig kapp	Ig kappa
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Db RESULT S41374 S41374 S41374 SACCE C; Spec C; Date C; Acces R; Arts R; Arts R; Desc.	Db Db	Que Bes Mai	RESULT A56446 C; pea C; Date C; Date C; Acce C; Acce A; Titl A; Titl A; Refe A; Refe A; Stat A; Role A; Resi A; Resi A; C; Keyw
Qy 240 FGGGTKLEIK 249	1	atch 65.2%; Score 857; DB 2; Length 268; cal Similarity 66.8%; Pred. No. 3.1e-54; 167; Conservative 26; Mismatches 45; Indels 12; Gs OVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWYKQRPEEGLEWIGYIDPESGETEY	RESULT 1 A56446 A56446 If heavy chain V region (3H-3H scFv) - mouse (strain BALB/C) C; Species: Mus musculus (house mouse) C; Date: 19-Jan-1996 #sequence_revision 19-Jan-1996 #text_change 16-Aug-1996 C; Accession: A56446 R; Tang, P.M.; Foltz, L.A.; Mahoney, W.C.; Schueler, P.A. J. Biol. Chem. 270; 7829-7835, 195 A; Title: A high affinity digoxin-binding protein displayed on M13 is functionally ide A; Reference number: A56446; MUID:95229583 A; Accession: A56446; MUID:95229583 A; Scatus: preliminary A; Molecule type: mRNA A; Residues: 1-568 <tan> A; Cross-references: 3B:U20617 C; Keywords: heterotetramer; immunoglobulin</tan>

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A; Reference number: S41374
A; Accession: S41374
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-249 < ART>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Jannot, C.B.; Hynes, N.E.
Biochem. Biophys. Res. Commun. 230, 242-246, 1997
A;Title: Characterization of scFv-421, a single-chain A;Reference number: JC5322; MUID:97168950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C:Species: Homo sapiens (man)
C:Date: 15-May-1997 #sequence_revision 15-May-1997
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A; Residues: 1-233 <JAN>
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Best Local S
Matches 165
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Best Local Similarity 68.3
Matches 172; Conservative
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                               245 KLEI 248
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 230 KLEI
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                                                                              QSPKVLIYWASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLPTFGGGT 244
                                                                                                                                         GGSGGGGSGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQQKPG
                                                                                                                                                                                                         QGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTVSSGG
                                                                                                                                                                                                                                                                    QESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEYAPNF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TVSSGGGGSGGGGGGGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAW 178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QVKLQESGAELVRSGASVNLSCTASGFNIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
                                                              QPPRLLIYLVSNLESGVPARFSGSGSGTDFTLNIHPVEEEDAATYYCQHIRELTRSEGGT
                                                                                                                          GGSGGRASGGGGSDIELTQSPASLAVSLGQRATISCRASKSV--STSGYSYMHWNQQKPG
                                                                                                                                                                                                                                                     QESGAELVRSGASVKLSCTTSGFNINDYYMHWVKKRPEQGLEWIGRIDPENGDADMTRSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TVSSRGGGSGGGGGGGGDIELTQSPPSVVVIPGESVSISCRSSKSLLYS-DGDSYLFW
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                                                                                                                                                                                                                                                                                                                  165;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FLQRPGQSPQLLIYRMSNLASGVPDRFSGSGSGTSFTLRISRVEAEDVGVYYCMQHREYP
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Pred. No. 3.4e-53;
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Ig kappa chain V.region (anti-DNA, DP13VK and DP18VK) - mouse (fragment) C:Species: Mus musculus (house mouse) C:Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000 C:Accession: PL0265
                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA'
A; Residues: 1-113 <BH>
A; Residues: 1-113 <BH>
C; Superfamily: immunoglobulin V region; immunoglobulin
C; Keywords: immunoglobulin
C; Keywords: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Accession: PT0407
R;Behar, S.M.; Lustgarten, D.L.; Corbet, S.; Scharff, M.D.
J. Exp. Med. 173, 731-741, 1991
A;Title: Characterization of somatically mutated S107 VH11
A;Reference number: FT0376; MUID:91147903
A;Accession: PT0407
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: S43103
R;Gilbert, D.; Brard, F.; Margaritte, C.; Delpech, A.; Tron, F.
submitted to the EMBI. Data Library, March 1994
A;Description: An id:otype D23-bearing polyspecific, murine anti-DNA monoclonal antib
A;Reference number: S42484
A;Accession: S43103
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A; Residues: 1-112 <GIL>
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C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change
  R; Shlomchik, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Status: preliminary
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Best Local S
Matches 104
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Best Local S
Matches 105
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Pred. No. 6.2e-32;
3; Mismatches 5
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Pred. No. 2.3e-32;
3; Mismatches 4
  Η.;
  Radic,
  M. Z.;
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Pisetsky,
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A; Residues:
C; Superfamil
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A;Title: Cloning and characterization of cDNAs coding for the heavy and light A;Reference number: PS0023; MUID:89232725
IT0408
Ig light chain V region (S107/VH11 group 1-6) - mouse (fragment)
C:Species: Mus musculus (house mouse)
C:Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change
C:Accession: PT0408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Note: the amino-terminal four residues of the mature protein were directly sequenced C;Comment: This chain was obtained from a monoclonal antibody against Pseudomonas aerug C;Superfamily: immunoglobulin v region; immunoglobulin homology C;Reywords: heterotetramer; immunoglobulin homology C;Reywords: signal sequence #status predicted <SIG>F;1-20/Domain: signal sequence #status predicted <SIG>F;21-133/Product: Ig kappa chain V region 6A4 #status experimental <IGV>F;36-116/Domain: immunoglobulin homology <IMM>
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A; Title: Anti-DNA a
A; Reference number:
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A; Molecule type: mRNA
A; Residues: 1-133 <MAR>
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F;63-94/Region: framework 3
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                                                                                                            RESULT
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Best Local
                                                                                                                                                                                                                                                                                                                                                         Matches 104;
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Best Local
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Title: Anti-DNA antibodies
                                                                                                                                                                                                      194 ASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLPTFGGGTKLEIK 249
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                         #sequence_revision 30-Sep-1993 #text_change 21-Jan-2000
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92.0%;
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89.7%;
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Pred. No. 1.7e-31;
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Pred. No. 1.4e-31;
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A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-112 <MAR>
A;Cross-references: EMBL:Z29536
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
                                                                                                                           Ig kappa chain V region (6188) - mouse (fragment) (Species: Mus musculus (house mouse) (C; Species: Pus musculus (house mouse) (C; Sate: 30-Sep-1991 #sequence_revision 30-Sep-1991 (C; Accession: $03304; JLD043 R; Van Cleave, V.H.; Naeve, C.W.; Metzger, D.W. J. Exp. Med. 157#;1841-1848, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig kappa chain V region (12.5H VL) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change
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J. Exp. Med. 173,
A;Title: Characte
A; Molecule type: mRNA
A; Residues: 1-111 <VAN>
A; Cross-references: EMBL:X12380
A; Note: the authors translated
                                                                        A;Title: Do antibodies recognize amino acid
A;Reference number: JL0043; MUID:88258372
A;Accession: S03304
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A; Accession: $41393
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A; Residues: 1-113 <BEH>
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A; Accession: PT0408
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Best Local Similarity
Matches 99; Conserv
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Best Local Similarity
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Characterization of somatically
                                                                                                                                                                                                                                                                                                                        DSGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLRTFGGGTKLEIK
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EMBL:X12380; NID:g52341; translated the codon GAT
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Pred. No. 1.2e-30;
3; Mismatches 8
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PIDN:CAA30938.1; I for residue 76 as
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Ig heavy chain precursor V region (MRL-histone 7H) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 04-Dec-1992 #sequence_revision 04-Dec-1992 #text_change 21-Jan-2000
C;Accession: S04576
R;Kofler, R.; Noonan, D.J.; Strohal, R.; Balderas, R.S.; Moller, N.P.H.; Dixon, F.J
Eur. J. Immunol. 17, 91-95, 1837
A;Title: Molecular analysis of the murine lupus-associated anti-self response: invo. A;Reference number: S04573; MUID:87133856
A;Recession: S04576
A;Molecule type: mRNA
A;Residues: 1-136 <a href="KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND">KCND"<KCND">KCND">KCND"<KCND">KCND">KCND"<KCND">KCND"<KCND">KCND">KCND"<KCND">KCND"<KCND">KCND"
                                                                                 Ig light chain V region (clone 111.19) - mouse C; Species; Mus musculus (house mouse) C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change C; Accession: PHI101

R; Tillman, D.M.; Jou, N.T.; Hill, R.J.; Marion, T.N.

J. Exp. Med. 176, 761.779, 1992

A; Title: Both IgM and IgG anti-DNA antibodies are the products of A; Reference number: PH0971; MUID:92381444

A; Accession: PHI101

A; Status: nucleic acid sequence not shown

A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-104 <TIL>
A; Experimental source: B cell, C; Superfamily: immunoglobulin C; Keywords immunoglobulin
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PH1101
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Matches 94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                          strain [NZB x NZW]F1
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Pred. No. 1.7e-29
4; Mismatches
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                             immunoglobulin
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A;Title: Both IgM and IgG anti-DNA antibodies are A;Reference number: FH0971; MUID:92381444
A;Accession: PH1102
                                                                                                                                                                                   A;Experimental source: B cell, strain [NZB x NZW]F1 C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                           A;Status: nucleic acid sequence not shown A;Molecule type: mRNA A;Residues: 1-104 <TIL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig light chain V region (clone 111.61) - mouse
C; Species: Mus musculus (house mouse)
C; Date: 30-Sep-1993 #sequence_revision 30-Sep-
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Score 497; DB 2; Pred. No. 4.7e-29; 4; Mismatches 5

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Gaps

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Length 104

immunoglobulin homology

37.8%; 91.3%;

PH1102

30-Sep-1993 #text_change 21-Jan-2000

are the products

of clonally selective

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C; Accession: PH1047; PH1049

R; Tillman, D.M.; Jou, N.T.; Hill, R.J.; Marion, T.N.

J. Exp. Med. 176, 761-779, 1992

A; Tille: Both IgM and IgG anti-UNA antibodies are the A; Reference number: PH0971; MUID:92381444

A; Accession: PH1047

A; Status: nucleic acid sequence not shown

A; Molecule type: mRNA
A; Residues: 1-103 <TIL>
A; Experimental source: B cell, strain [NZB x NZW]F1
A; Experimental source: B cell, strain [NZB x NZW]F1
C; Superfamily: immunoglobulin
C; Keywords: immunoglobulin
C; Keywords: immunoglobulin homology <IMM>
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Pred. No. 4.6e-29;
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
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J. Exp. Med. 176, 761-779, 192
A;Title: Both IgM and IgG anti-DNA antibodies are the products of clonally selective B A;Reference number: PH0971; MUID:92381444
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C;Species: Mus musculus (house mouse)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 21-Jan-2000
C;Accession: PH1051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Accession: PH1051
A;Status: nucleic acid sequence not shown
A;Mclecule type: mRNA
A;Residues: 1-103 <TIL>
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Best Local Similarity 92.3
Matches 95; Conservative
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Copyright (c) 1993 - 2000 Compugen Ltd
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Query Match, 35.3%; Best Local Similarity 75.0%; Matches 87; Conservative 1

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Length 133;

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Gaps

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01-APR-1988 (Rel. 07, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
IG KAPPA CHAIN V-IV REGION B17 PRECURSOR.
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MEDLINE-86041854; PubMed-2997713;

Marsh P., Mills F., Gould H.;

"Detection of a unique human V kappa
cDNA probe.";
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COMPLEMENTARITY-DETERMINING
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RESULT 4
KV40_HUMAN
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KV40_HUMAN STANDER...

P06312;

01-JAN-1988 (Rel. 06, Created)

01-JAN-1988 (Rel. 06, Last sequence update)

15-JUL-1999 (Rel. 38, Last annotation update)

15-JUL-1999 (Rel. 38, Last annotation update)

15-JUL-1999 (Rel. 38, Last annotation update)
                   "Subgroup IV of human immunogiodulii n 119...
Single germline gene.";
Nucleic Acids Res. 13:6515-6529(1985).
Nucleic MISCELLANEOUS: THERE IS ONLY ONE IG KAPPA V-IV GENE.
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DISULFID
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SEQUENCE
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This SWISS-PROT entry between the Swiss Ins
                                                                                 Zachau H.G.
                                                                                            Klobeck H.G.,
                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; ; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted (AUG-1996) to -!- MISCELLANEOUS: THE C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "The primary structure of a monoclonic immunoglobulin-L-chain subgroup IV of the kappa type (Bence-Jones protein Len.)."; hoppe-Seyler's Z. Physiol. Chem. 356:507-557(1975).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-1986 (Rei. 01, Created)
01-CCT-1996 (Rei. 34, Last sequence
15-JUL-1999 (Rei. 38, Last annotatio
IG KAPPA CHAIN V-IV REGION LEN.
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HSSP; P01789; 2MCP.
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                                                                                                        MEDLINE=86041853;
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86; Conserv
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schmann N.;
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REGION OF THIS CHAIN HAS
IS A BENCE-JONES PROTEIN.
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Pred. No. 1.4e-26;
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COMPLEMENTARITY-DETERMINING
FRAMEWORK 4.
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 jht. It is produced through a collaboration
Bioinformatics and the EMBL outstation -
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European Bioinformatics Institute.

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                                                                                                                         P01751; P01752;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
IG HEAVY CHAIN V REGION B1-8/186-2 PRE
Mus musculus (Mouse).
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                                                                               This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no rest
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SIGNAL
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Bothwell A.L.M., Paskind M., Reth M.,
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                    EMBL;
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                    J00529; AAA38170.1;
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Rodentia;
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Pred. No. 2
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COMPLEMENTARITY-DETERMINING
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COMPLEMENTARITY-DETERMINING
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FRAMEWORK 1
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RESULT 6
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                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE-81241357; Pubmed-6788890;
Kwan S.-P., Rudikoff S., Seidman
                                                                                                                                                                                                                                                                                      Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                        21-JUL-1986
15-JUL-1999
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                                                                                                          EMBL; U29423;
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                                                                                                                                                                                                              chains.
                                                                                                                                                                                                                                                                    Mammalia;
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Rodentia;
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Seidman J.G., I
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Pred. No. 6
                           FRAMEWORK 1.

COMPLEMENTARITY - DETERMINING
FDAMEWORK 2.

TOTAL PROPERTINING
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COMPLEMENTARITY-DETERMINING
FRAMEWORK 3.
D SEGMENT.
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COMPLEMENTARITY-DETERMINING
                          FRAMEWORK 2.
COMPLEMENTARITY-DETERMINING
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P01633;

`21-JUL-1986

21-JUL-1986

15-JUL-1999
                                                                                                                                                                                                                                                                                      INTERPRO; IPRO33006; -.
PFAM; PF00047; 19; 1.
Immunoglobulin V region; Antiarsonate antibody; Hybridoma
NON TER 120 120
SEQUENCE 120 AA; 13207 .... ---
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NON_TER
SEQUENCE
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Eur. J. Immunol. 12:1023-1032(1982).
-i- MISCELLANEOUS: FROM ANALYSIS OF THE SIZES OF SEVERAL
DIFFERENTIATED GENES THAT HYBRIDIZE TO THIS ONE, THE
CONCLUDE THAT ALL OF THESE V REGIONS HAVE REARRANGED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P01747;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1996 (Rel. 01, Last seq
15-JUL-1999 (Rel. 38, Last ann
IG HEAVY CHAIN V REGION 36-65.
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Eukaryota; Metazoa; (
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A02028; HVMSG7.
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                                                                                                                                                                    PNFQGKATVTADTSSNTAYLHLSSLTSEDTTVYYCNAVIYYEYDGYALDYWGQGTTVTVS 121
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Rodentia;
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Pred. No. 1.5e
20; Mismatches
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Pred. No. 1.4e
L6; Mismatches
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Sciurognathi; Muridae;
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n A mouse.";
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; Murinae; Mus
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Blockhem. J. 171:337-347(1978).

-i- MISCELLANEOUS: THE MATURE CHAIN HAS 12 ADDITIONAL RESIDUES AMINO END, DUE TO A TANDEM DUPLICATION OF 36 NUCLEOTIDES AF CODON FOR RESIDUE 36. RESIDUE 42 CORRESPONDS TO THE AMINO-T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kelley D.E., Colectough C., Perry K.F.; "Functional significance and evolutionary development 5'-terminal regions of immunoglobulin variable-region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE OF 1-71 FROM N.A.
MEDLINE-83001944; PubMed-6288267;
Kelley D.E., Colectough C., Perry R.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Meta:oa;
Mammalia; Eutheria;
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MEDLINE=78186617; PubMed=418775;
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"The variability, arrangement, and
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                                                                          134
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                                                                                                                                                                                                                                                                                                                                 [mmunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Sequence of the full-length immunoglobulin kappa-chain
                       194 ASTRESGYPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNL-PTFGGGTKLEIK
92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESIDUE OF TYPICAL KAPPA CHAINS.
                                                               GGGSDIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQQKPGQSPKVLIYW 193
                                                                                                                                                                                                                                                                                                                                                                      A01916; KVMS11
ASYRYTGVPDRFTGSGSGTDFTFTISSVQAEDLAVYYCQQHYSTPPTFGGGTKLEIK
                                                GVDGDIVMTQSHKFMSTSVGDRVSITCKASQDVSTT-----VAWYQQKPGQSPKLLIYS 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Biochem, 58:176-187(1980).
                                                                                                7,6;
                                                                                                               Similarity
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                                                                                                   Conservative
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Pred. No. 1.
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COMPLEMENTARITY-DETERMINING 2.
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.7e-21;
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21-JUL-1986
21-JUL-1986
15-JUL-1999
IG HEAVI CHALL.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Eukaryota; Fintheria; Rodentia;
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Mus musculus (Mouse).

Mus musculus (Mondata;

Mortazoa; Chordata;

Mortazoa; Rodentia;
                                                                P01755;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1996 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
IG HEAVY CHAIN V REGION S43 PRECURSOR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "The complete amino acid sequence of a mouse kappa light chain."; Biochem. J. 128:427-444(1972).
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MEDLINE=73053310; PubMed=4638343;
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                                                                                                                                                                                                                                                                                         ASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLF-TFGGGTKLEIK 249
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                                                                                                                                                                                                                                                                                                                                                 GADGNIVMTQSPKSMSMSVGERVTLTCKASENVV-----TYVSWYQQKPEQSPKLLIYG 79
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(Rel. 38, Mary Precursor.
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Pred. No. 2.2e
18; Mismatches
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COMPLEMENTARITY-DETERMINING
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COMPLEMENTARITY-DETERMINING
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        Craniata; Vertebrata; I
Sciurognathi; Muridae;
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                            Euteleostomi;
          Murinae;
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                                                                                HV02_MOUSE STANDARD; PRT; 140 AM.
P01746;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
16 HEAVY CHAIN V REGION 93G7 PRECURSOR.
                                                                                                                                                MOUSE
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SEQUENCE
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DOMAIN
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DOMAIN
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Cell 24:625-637(198)
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            MEDLINE=82152818;
Sims J., Rabbitts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Heavy chain variable region antibodies: somatic mutation
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SEQUENCE FROM
                               STRAIN-A,
                                        SEQUENCE FROM N'A.
                                                                               Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the European Bioinformatics Institute.
                                                            Mammalia; Eutheria;
                                                                     Eukaryota; .Metazoa;
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            PubMed=6801765;
T.H., Estess P.,
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15200
                                                           Chordata;
Rodentia;
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61.5%;
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JH2 SEGMENT.
BY SIMILARITY
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FRAMEWORK 1.
COMPLEMENTARITY-DETERMINING
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                                                           Sciurognathi;
                                                                     Craniata; Vertebrata; Euteleostomi;
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SO THE WAR AREA COCCOCCE

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Matches
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Best Local
                                                                                           Zakut R., Cohen J.,
Nucleic Acids Res.
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21-JUL-1986
21-JUL-1986
15-JUL-1999
                                                                                                                                                            SEQUENCE FROM N.A.

MEDLINE-81053741; PubMed-6253904;

Zakut R., Cohen J., Givol D.;

"Cloning and sequence of the cDNA corresponding region of immunoglobulin heavy chain MPC11.";

Nucleic Acids Res. 8:3591-3601(1980).
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SEQUENCE
                                                                                                                                                                                                                                                                                   21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
IG HEAVY CHAIN V REGION MPC 11.
Mus musculus (Mouse).
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PIR; 1
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            NON
                         PFAM; PF00047; ig; 1.
Immunoglobulin V region.
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PFAM; PF00047; ig; 1.
Immunoglobulin V region; Antiarsonate antibody; Hybridoma;
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                                                                               leic Acids Res. 8:4839-4840(1980).
MISCELLANEOUS: THIS SEQUENCE WAS TRANSLATED
FROM A MYELOMA THAT SECRETES IGG2B.
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0047; ig; 1.
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nes 31;
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                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Klobeck H.G., Combriato G., Zachau H.G.; "Immunoglobulin genes of the kappa light c lymphoid cell lines are closely related.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HUMAN
                                                                                                                                                                                                                               DOMAIN
                                                                                                                                                                                                                                                     SIGNAL
                                                                                                                                                                                                                                                                                                      EMBL; X00965; CAA25477.1; PIR; A01883; K1HUWK.
                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic
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            194
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 QVKLQESGAELVRSGASVNLSCTASGENIKDYYMHWVKQRPEEGLEWIGYIDPESGETEY
                                            GGGSDTELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQQKPGQSPKVLIYW 193
            ASTRESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYN-LPTFGGGTKLEIK
                                GARCDIQMTQSPSSLSASVGDRVTITCRASQSI-----SNYLNWYQQKPGKAPKLLIYA
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P01607;
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| 72; | Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                          Acids Res. 12:6995-7006(1984).
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IPR003006;
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                                                                                                                                                           IG KAPPA CHAIN V-I REGION WALL FRAMEWORK 1.
COMPLEMENTARITY-DETERMINING 1 FRAMEWORK 2.
COMPLEMENTARITY-DETERMINING 2 FRAMEWORK 3.
COMPLEMENTARITY-DETERMINING 3 FRAMEWORK 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
                                                                           20;
                                                                                     Score 371.5; DB 1 Pred. No. 6.6e-21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi;
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                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    precursor.
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RESULT 14
KV1M_HUMAN
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KV1H_HUMAN
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Best Local Similarity
Matches 71; Conser
                                                                                         P01500;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                         DOMAIN
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 Watanabe S., Hilschmann N.;
"The primary structure of a
                                                         Homo sapiens (Human)
Eukaryota; Metazoa; (
Mammalia; Eutheria; )
                                                                                                                                                        KV1H_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                    DOMAIN
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HSSP; P01607; 1REI.
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15-JUL-1999 (Rel. 38, Last ann
IG KAPPA CHAIN V-I REGION LAY.
                       SEQUENCE.
MEDLINE=71032830; PubMed=4097974;
                                                                                                                                                                                                                                                                                                                                                                                                        DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IgM anti-gamma globulins (Lay/Pom) with shared
specificities.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IND. J. IMMUNO1. 5:677-684(1976).
MISCELLANBOUS: THE SECOND AND THIRD HYPERVARIABLE REGIONS OF THIS CHAIN ARE IDENTICAL WITH THOSE OF THE HUMAN POM V-III KAPPA CHAIN, WITH WHICH IT SHARES CERRAIN IDIOTYPIC DETERMINANTS.
MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM AN IGM WITH ANTI-GAMMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GLOBULIN ACTIVITY.
                                                                                                                                                                                                                             ESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNL-PTFGGGTKLEIK
                                                                                                                                                                                                                                                                          DIELTQSPSSLAVSAGEKVAMSCKSSQSLFNSRTRKNYLAWYQQKPGQSPKVLIYWASTR 197
primary structure of a monoclonal kappa-type immunoglobulin
                                                                                                                                                                                                               EAGVPSRFSGSGSGTDFTFTISSLQPEDIATYYCQQYNNWPPTFGQGTKVEVK 107
                                                                                                                                                                                                                                                             DIQMTQSPSSLSVSVGDRVTITCQASQNV-----NAYLNWYQQKPGLAPKLLIYGASTR 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          J. Immunol.
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(Rel. 01, Last sequence update)
(Rel. 38, Last annotation update)
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                                                           Chordata;
Primates;
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Pred. No. 7.
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BY SIMILARITY.
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                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                        PRT;
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Job time: 195 sec
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Matches 70
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-!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE INV (3) MARKER
-!- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                 chain of subgroup I (Bence-Jones Protein Hau): subdivision within subgroups.";
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                                                 PSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQNYITPTSFGQGTRVEIK
                                                                       ESGVPDRFTGSGSGTDFTLTISSVQAEDLAVYYCKQSYNLPT-FGGGTKLEIK 249
                                                                                            DIQMTQSPSSLSASVGDRVTITCRASQSI-----SSYLSWYQQKPGKAPQVLIYAASSL
                                                                                                                DIELTOSPSSLAVSAGEKVAMSCKSSOSLFNSRTRKNYLAWYQQKPGQSPKVLIYWASTR 197
                                                                                                                                                                                                                                                                                                                               P80362;
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                                                                                                                                      Conservative
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         2001, 15:26:39
                                                                                                                                    Score 367.5; DB 1
Pred. No. 1.1e-20;
2; Mismatches 14
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COMPLEMENTARITY-DETERMINING FRAMEWORK 4.
BY SIMILARITY.
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